

pomegranate

fast and flexible probabilistic modelling in python

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UNIVERSITY *of* WASHINGTON

eScience Institute

ADVANCING DATA-INTENSIVE DISCOVERY IN ALL FIELDS

Inria
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Overview

pomegranate is **more flexible** than other packages, **faster**, is **intuitive to use**, and can do it all in **parallel**

Probability
Distributions

Bayes Classifiers

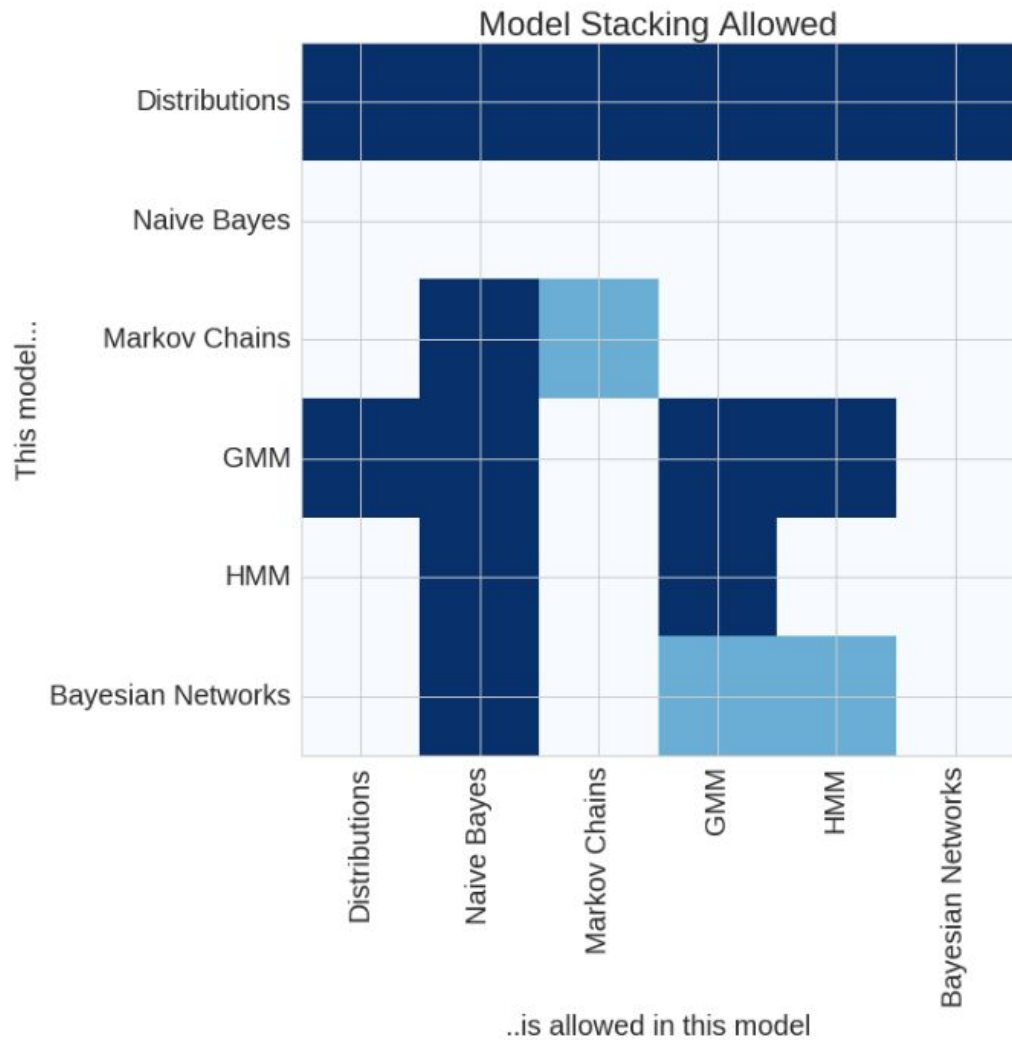
Bayesian
Networks

Markov Chains

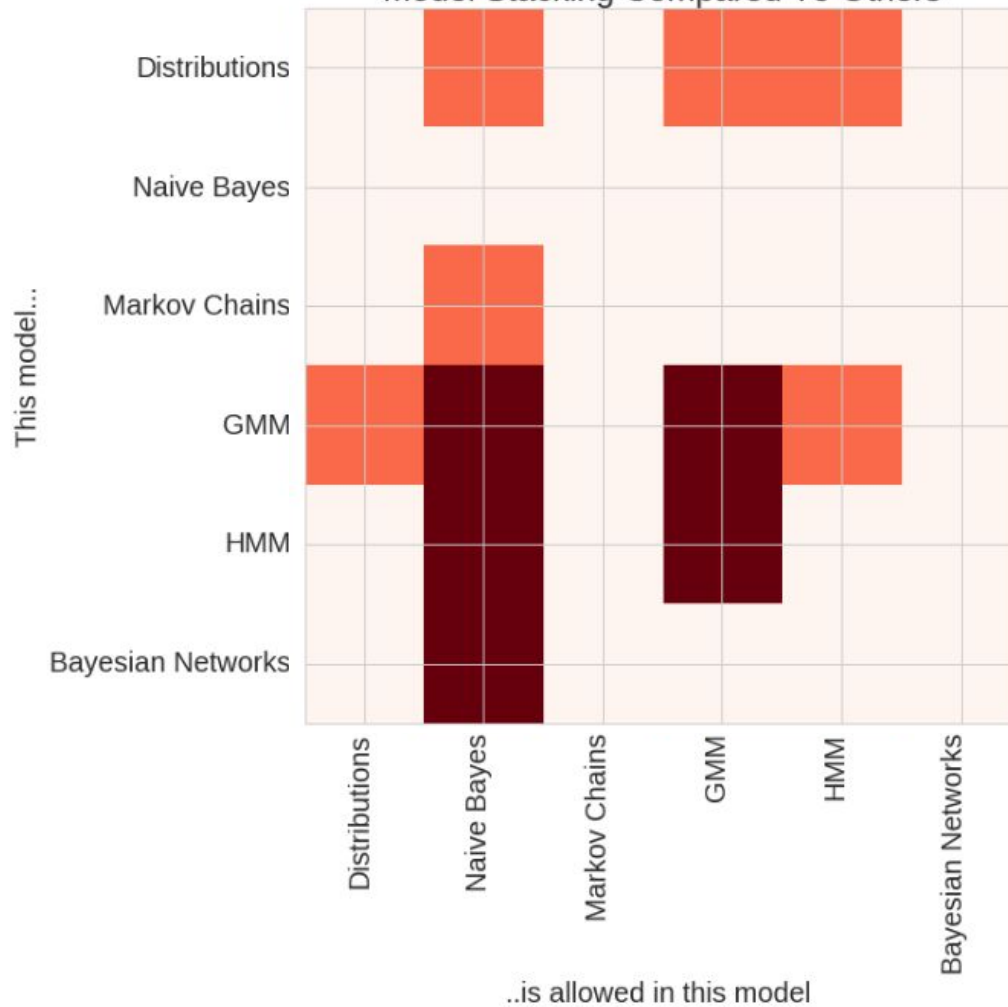
Hidden Markov
Models

General Mixture
Models





Model Stacking Compared To Others



Overview

The API

Major Models/Model Stacks

1. General Mixture Models
2. Hidden Markov Models
3. Bayesian Networks
4. Bayes Classifiers

Parallelization

Finale: Train a mixture of hidden markov models in parallel

All models share most methods

`model.log_probability(X) / model.probability(X)`

`model.sample()`

`model.fit(X, weights, inertia)`

All models have these methods!

`model.summarize(X, weights)`

`model.from_summaries(inertia)`

`model.predict(X)`

`model.predict_proba(X)`

`model.predict_log_proba(X)`

All models composed of distributions (like GMM, HMM...) have these methods too!

`Model.from_samples(X, weights)`

All models except HMMs have this (coming soon!)

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`model.log_probability(X) / model.probability(X)`

`model.sample()`

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All models composed of distributions (like GMM, HMM...) have these methods too!

`Model.from_samples(X, weights)`

All models except HMMs have this (coming soon!)

pomegranate supports many distributions

Univariate Distributions

1. UniformDistribution
2. BernoulliDistribution
3. NormalDistribution
4. LogNormalDistribution
5. ExponentialDistribution
6. BetaDistribution
7. GammaDistribution
8. DiscreteDistribution
9. PoissonDistribution

Kernel Densities

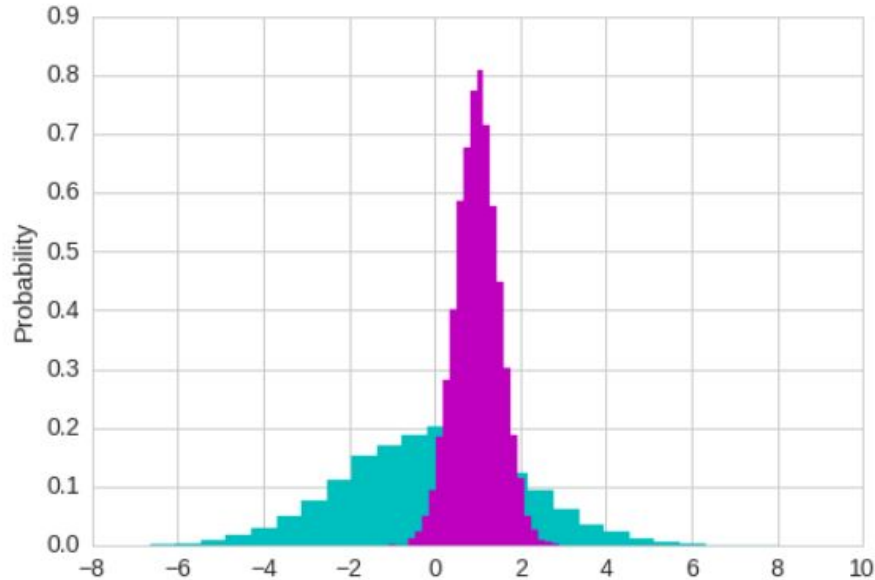
1. GaussianKernelDensity
2. UniformKernelDensity
3. TriangleKernelDensity

Multivariate Distributions

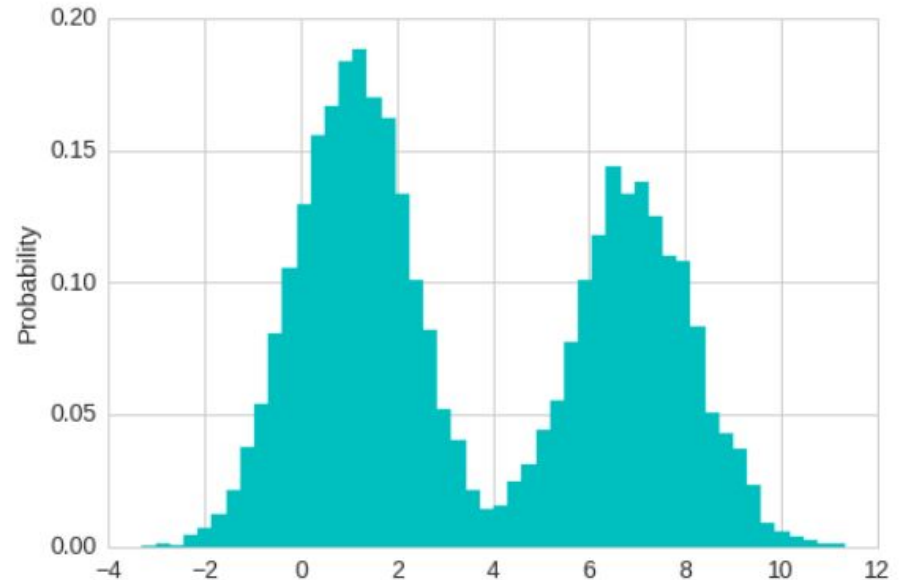
1. IndependentComponentsDistribution
2. MultivariateGaussianDistribution
3. DirichletDistribution
4. ConditionalProbabilityTable
5. JointProbabilityTable

Models can be created from known values

```
mu, sig = 0, 2  
a = NormalDistribution(mu, sig)
```

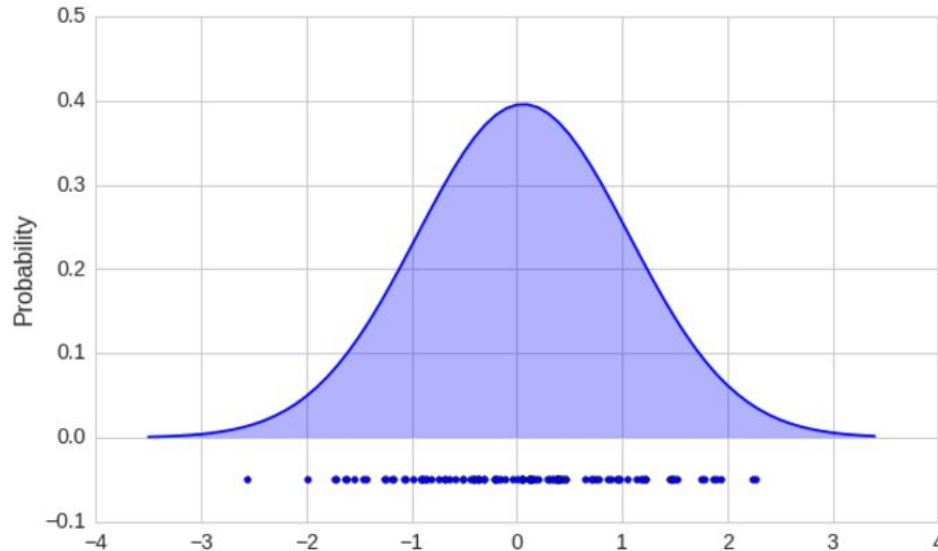


```
X = [0, 1, 1, 2, 1.5, 6, 7, 8, 7]  
a = GaussianKernelDensity(X)
```



Models can also be learned directly from data

```
X = numpy.random.normal(0, 1, 100)  
a = NormalDistribution.from_samples(X)
```



pomegranate can be faster than numpy

Fitting a Normal Distribution to 1,000 samples

```
data = numpy.random.randn(1000)

print "numpy time:"
%timeit -n 100 data.mean(), data.std()
print
print "pomegranate time:"
%timeit -n 100 NormalDistribution.from_samples(data)
```

numpy time:
100 loops, best of 3: 46.6 μ s per loop

pomegranate time:
100 loops, best of 3: 22.2 μ s per loop

pomegranate can be faster than numpy

Fitting Multivariate Gaussian to 10,000,000 samples of 10 dimensions

```
data = numpy.random.randn(10000000, 10)

print "numpy time:"
%timeit -n 10 data.mean(), numpy.cov(data.T)
print
print "pomegranate time:"
%timeit -n 10 MultivariateGaussianDistribution.from_samples(data)
```

numpy time:
10 loops, best of 3: 1.02 s per loop

pomegranate time:
10 loops, best of 3: 799 ms per loop

pomegranate can be faster than numpy

pomegranate reduces data to sufficient statistics for updates and so only has to go datasets once (for all models).

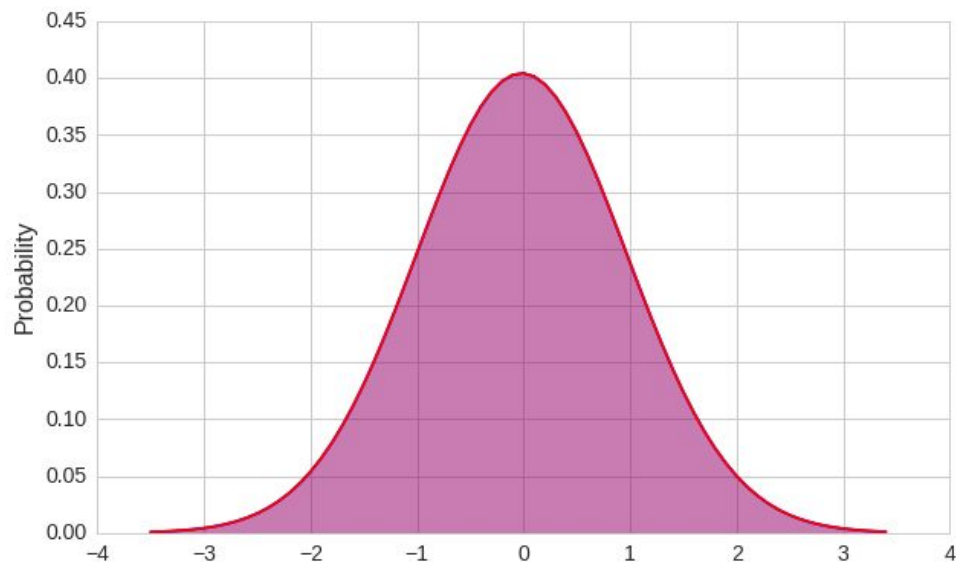
Here is an example of the Normal Distribution sufficient statistics

$$\sum_{i=1}^n w_i \quad \sum_{i=1}^n w_i x_i \quad \sum_{i=1}^n w_i x_i^2 \quad \longrightarrow \quad \begin{aligned} \mu &= \frac{\sum_{i=1}^n w_i x_i}{\sum_{i=1}^n w_i} \\ \sigma^2 &= \frac{\sum_{i=1}^n w_i x_i^2}{\sum_{i=1}^n w_i} - \frac{\left(\sum_{i=1}^n w_i x_i \right)^2}{\left(\sum_{i=1}^n w_i \right)^2} \end{aligned}$$

pomegranate supports out of core learning

Due to the use of sufficient statistics that are additive, pomegranate can natively support out-of-core/online learning, where you may not have the entire dataset in memory at a time

```
a.fit(data)
b.summarize(data[:1000])
b.summarize(data[1000:2000])
b.summarize(data[2000:3000])
b.summarize(data[3000:4000])
b.summarize(data[4000:])
b.from_summaries()
```



Fit Mean: -0.0174820965846, Fit STD: 0.986767322871
Summarize Mean: -0.0174820965846, Summarize STD: 0.986767322871

pomegranate can be faster than scipy

```
from scipy.stats import norm

d = NormalDistribution(0, 1)

print "scipy time:"
%timeit -n 100 norm.logpdf(2, 0, 1)
print
print "pomegranate time:"
%timeit -n 100 NormalDistribution(0, 1).log_probability(2)
print
print "pomegranate with (w/ created object)"
%timeit -n 100 d.log_probability(2)
print
print "logp difference: {}".format( norm.logpdf(2, 0, 1) - No
```

scipy time:
100 loops, best of 3: 96.3 μ s per loop

pomegranate time:
100 loops, best of 3: 560 ns per loop

pomegranate with (w/ created object)
100 loops, best of 3: 119 ns per loop

logp difference: -3.99236199655e-13

scipy: 96.3 us

pomegranate: 560 ns

pomegranate (w/ precreated object): 119 ns

pomegranate can be faster than scipy

pomegranate uses aggressive caching of values required for probability calculations to speed them up

$$P(X|\mu, \sigma) = \frac{1}{\sqrt{2\pi}\sigma} \exp\left(-\frac{(x - \mu)^2}{2\sigma^2}\right)$$

$$\log P(X|\mu, \sigma) = -\log(\sqrt{2\pi}\sigma) - \frac{(x - \mu)^2}{2\sigma^2}$$

$$\log P(X|\mu, \sigma) = \alpha - \frac{(x - \mu)^2}{\beta}$$



Example 'blast' from Gossip Girl

Spotted: Lonely Boy. Can't believe the love of his life has returned. If only she knew who he was. But everyone knows Serena. And everyone is talking. Wonder what Blair Waldorf thinks. Sure, they're BFF's, but we always thought Blair's boyfriend Nate had a thing for Serena.

Example 'blast' from Gossip Girl

Why'd she leave? Why'd she return? Send me all the deets.
And who am I? That's the secret I'll never tell. The only one.
—XOXO. Gossip Girl.

How do we encode these blasts?

Better lock it down with Nate, B. Clock's ticking.

+1 Nate

-1 Blair

How do we encode these blasts?

Better lock it down with Nate, B. Clock's ticking.

+1 Nate

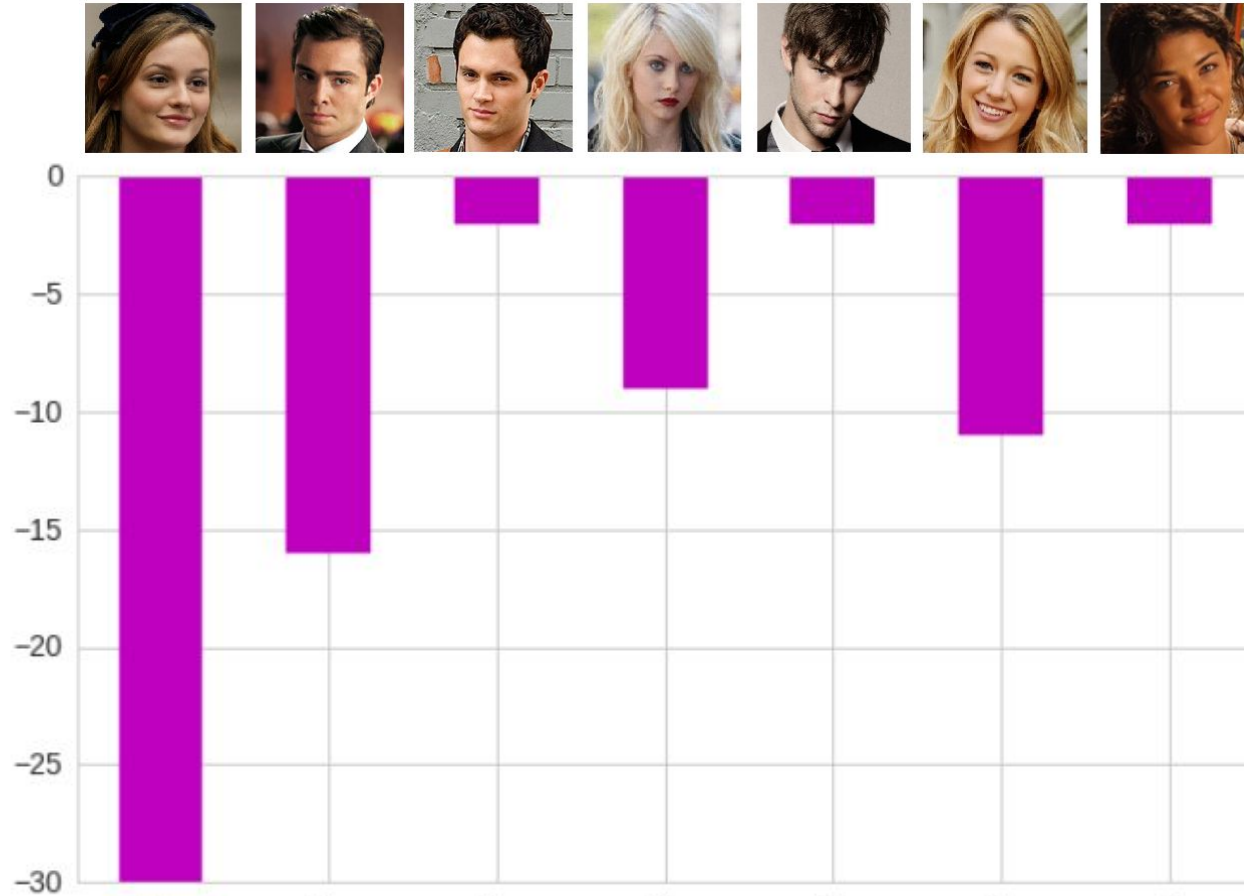
-1 Blair

This just in: S and B committing a crime of fashion. Who doesn't love a five-finger discount. Especially if it's the middle one.

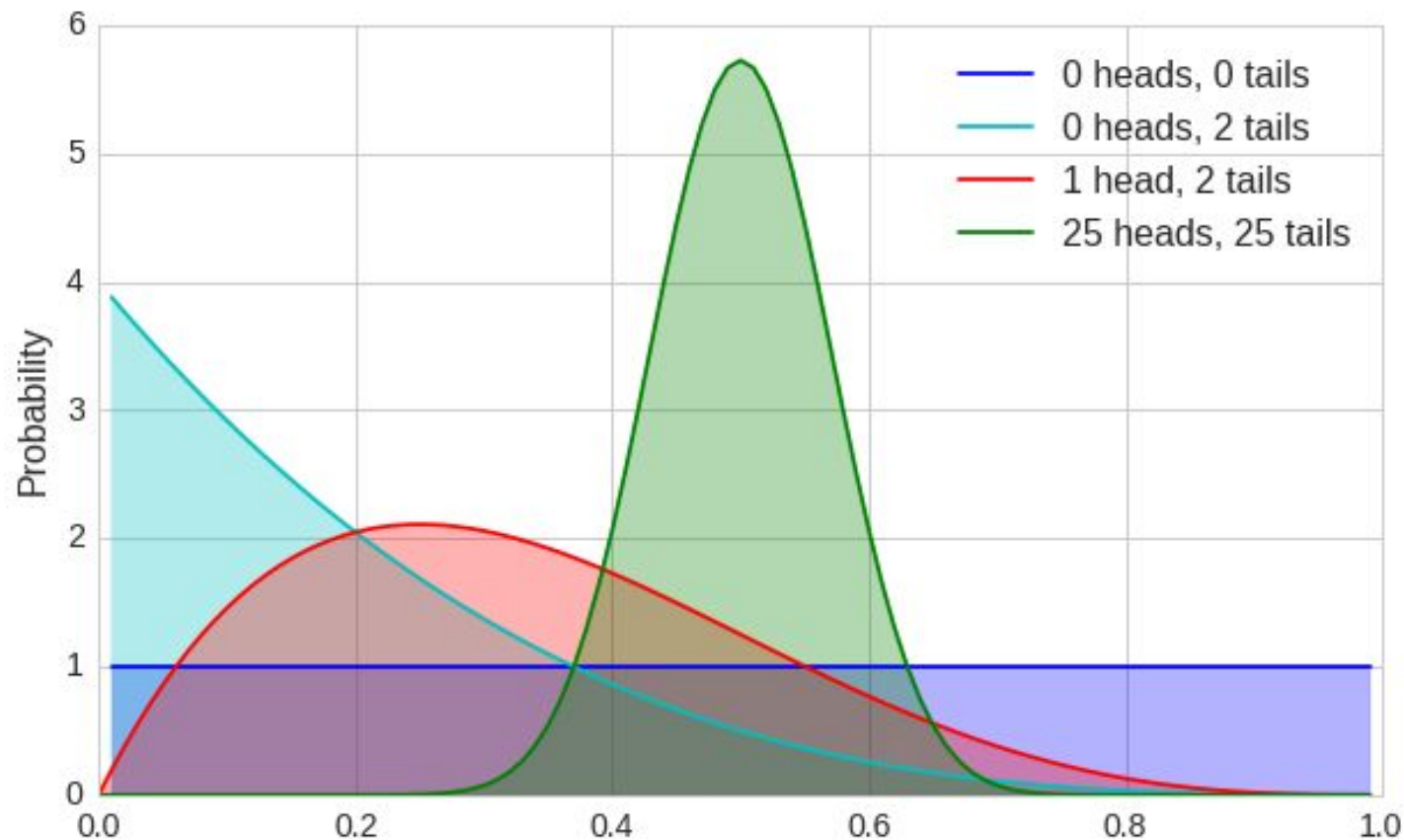
-1 Blair

-1 Serena

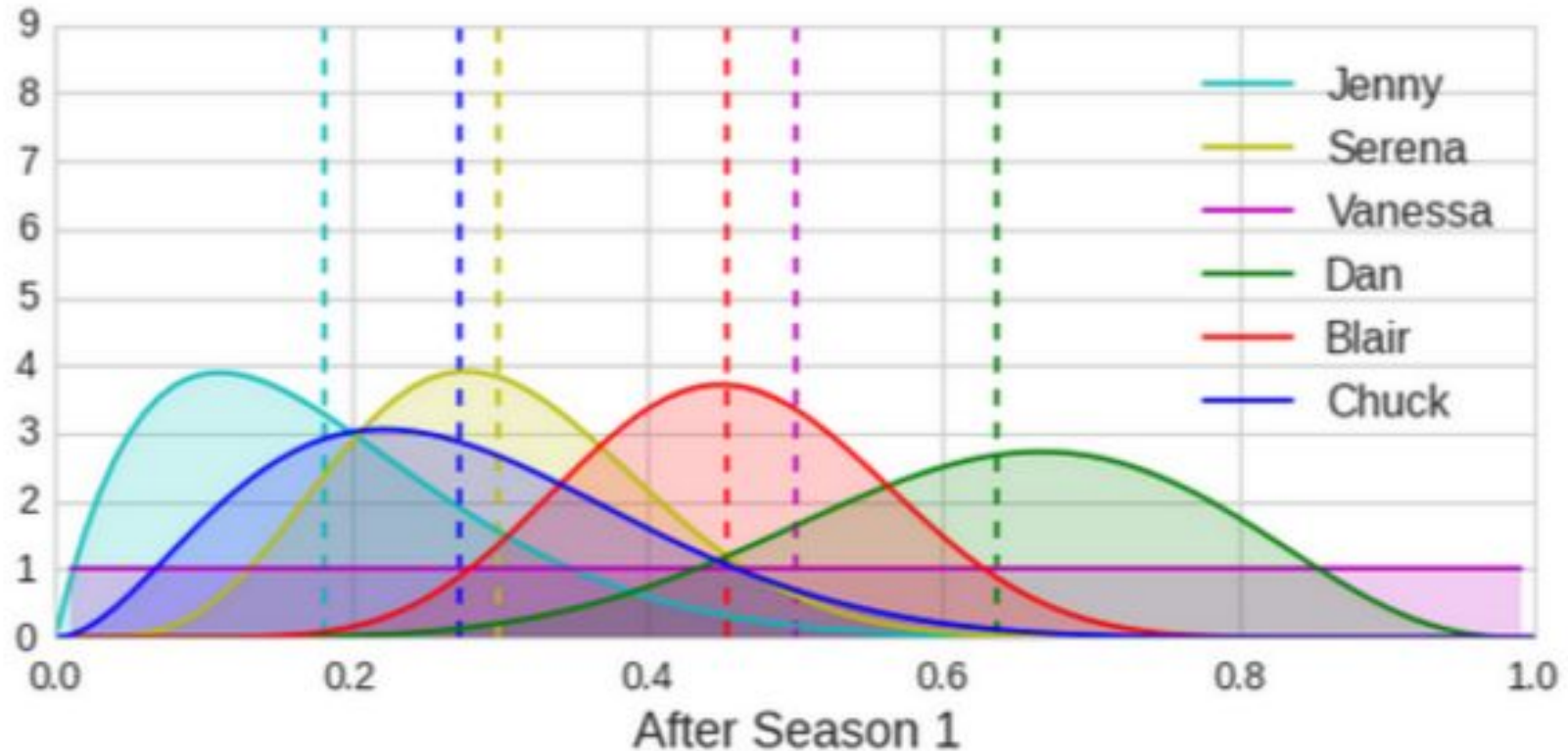
Simple summations don't distinguish well



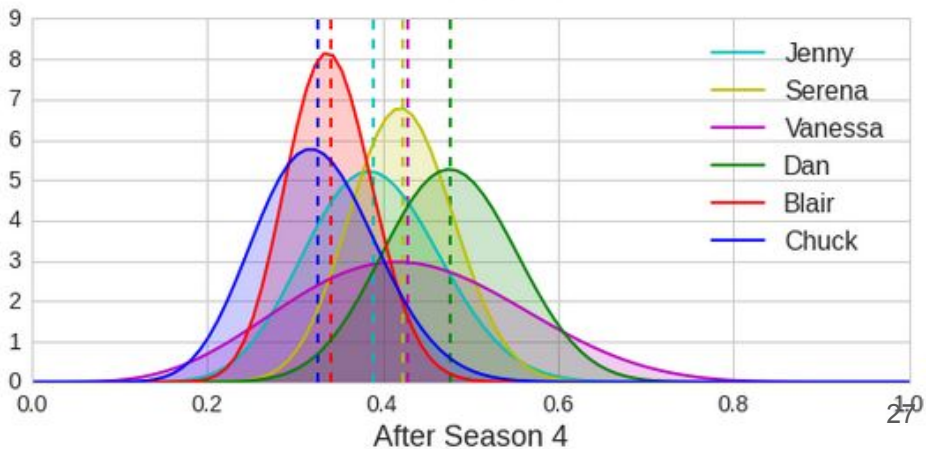
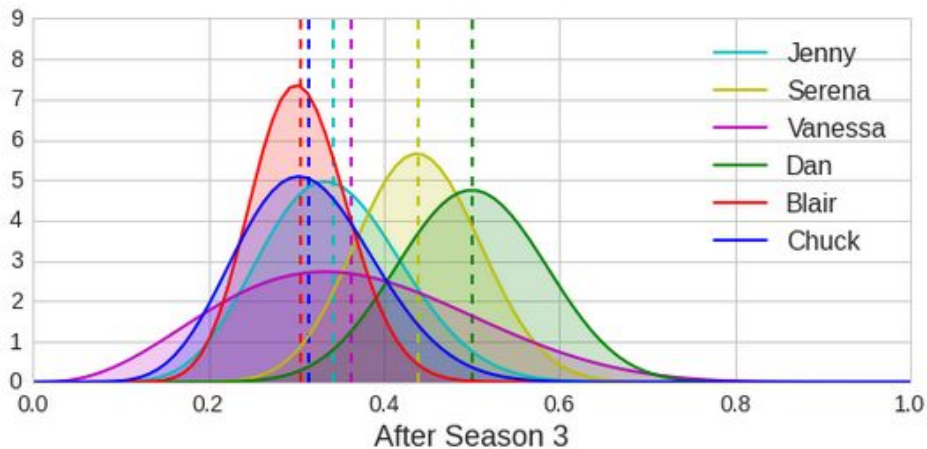
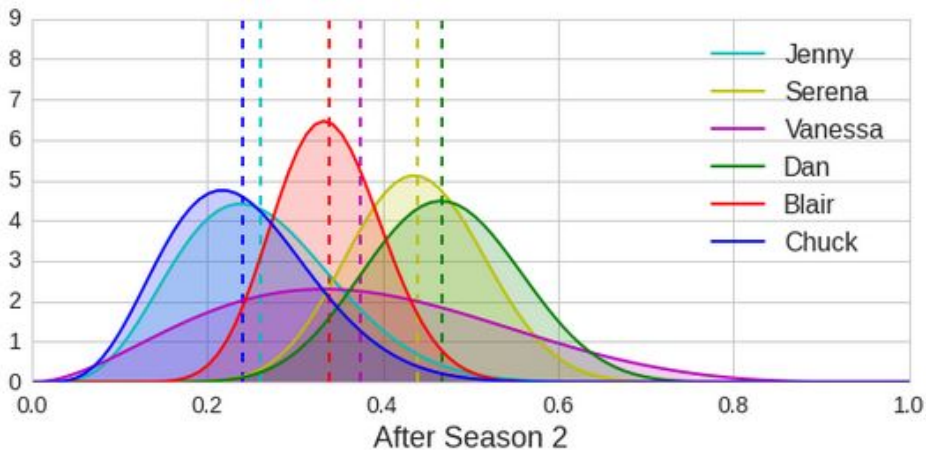
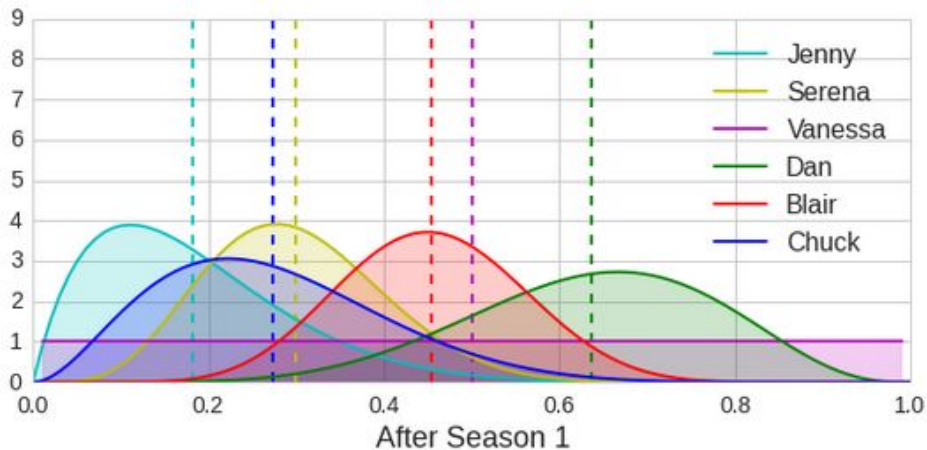
Beta distributions can model uncertainty well



Beta distributions capture our certainty about the identity of Gossip Girl



The distributions converge as the show progresses



Overview

The API

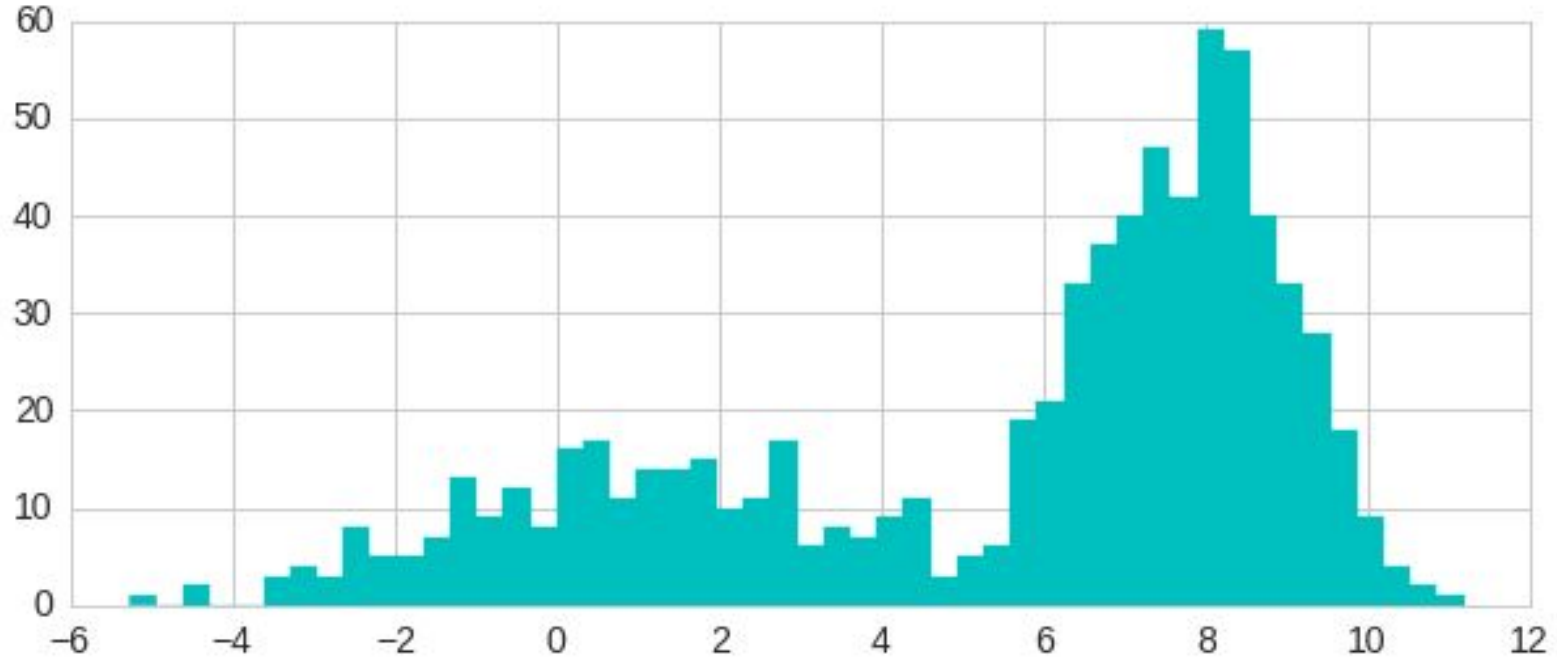
Major Models/Model Stacks

1. General Mixture Models
2. Hidden Markov Models
3. Bayesian Networks
4. Bayes Classifiers

Parallelization

Finale: Train a mixture of hidden markov models in parallel

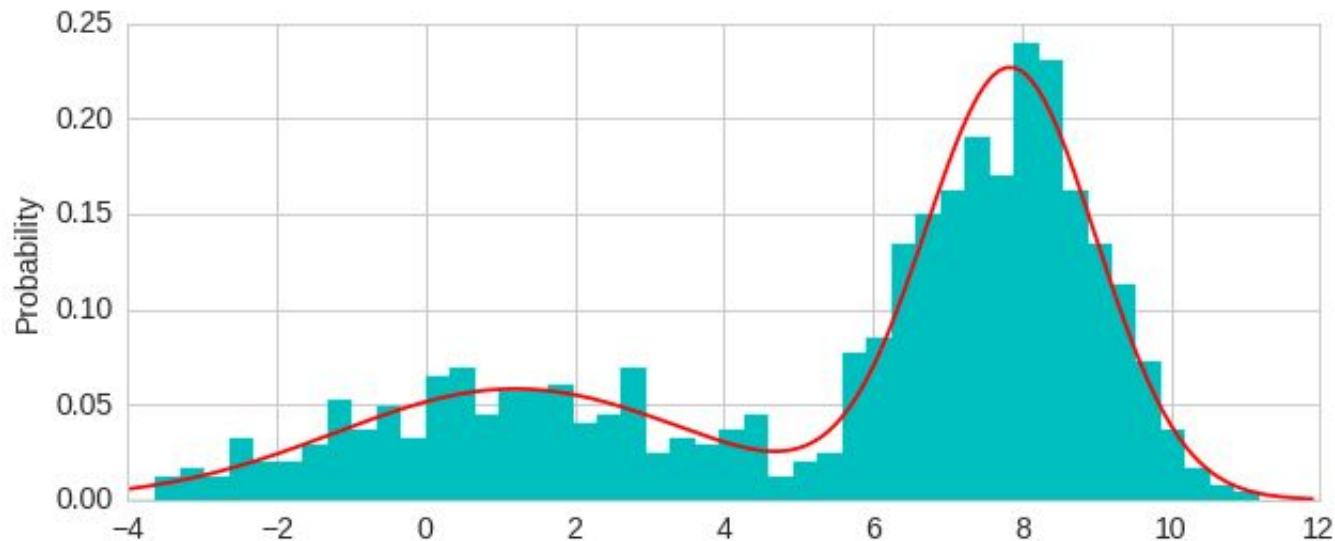
General Mixture Models (GMMs) can model multi-component distributions



GMMs use Expectation-Maximization (EM) to fit

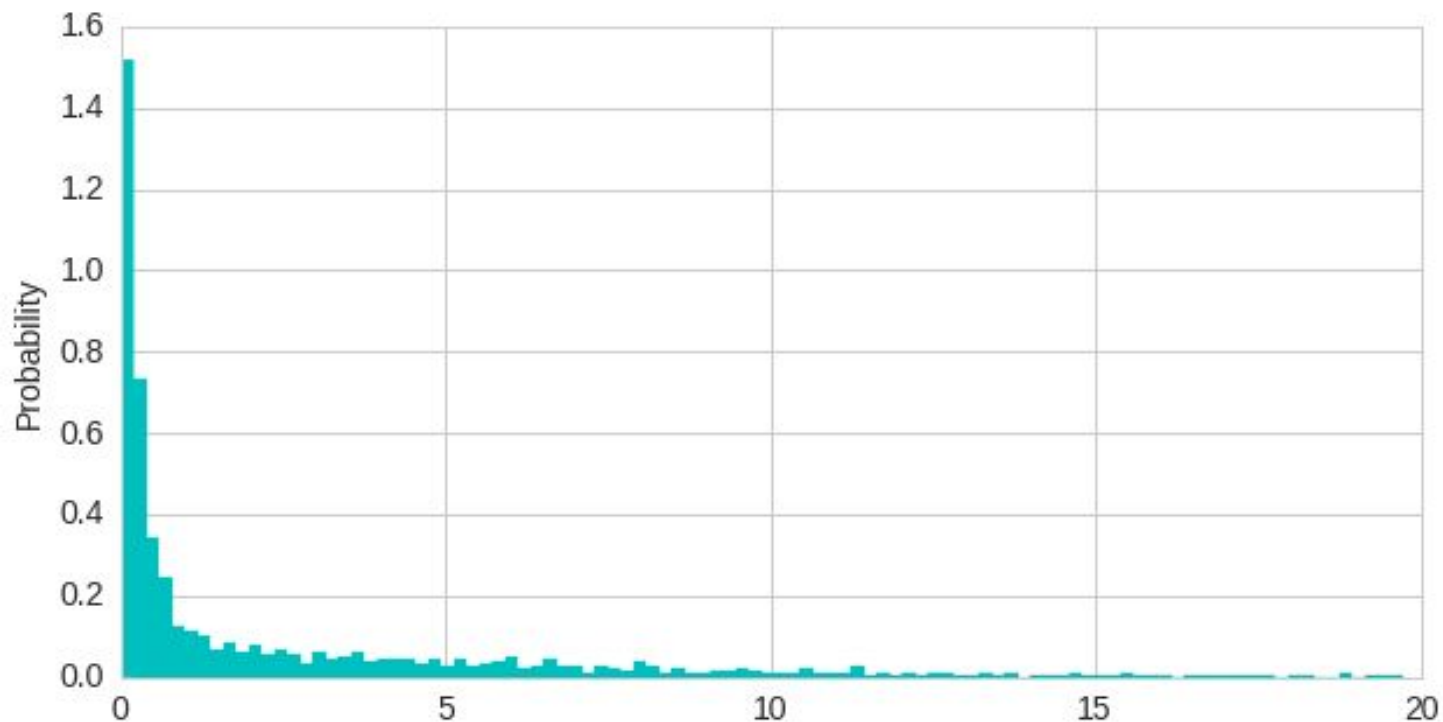
1. Initialize clusters using kmeans++ or kmeans||
2. Assign weights to all points equal to the posterior $P(M|D)$ (E step)
3. Update distribution using weighted points (M step)
4. Repeat 2 and 3 ~~forever~~ until convergence

General Mixture Models (GMMs) can model multi-component distributions

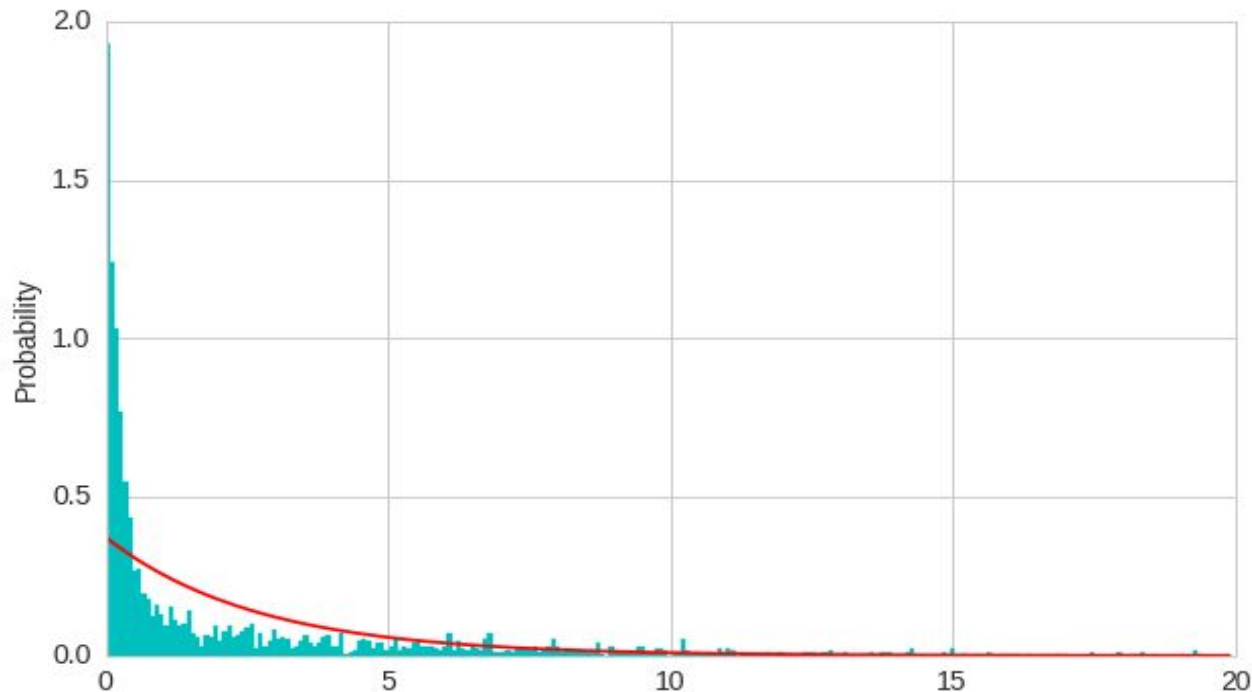


```
model = GeneralMixtureModel.from_samples(NormalDistribution, 2, X)
```

GMMs are not limited to Gaussian distributions

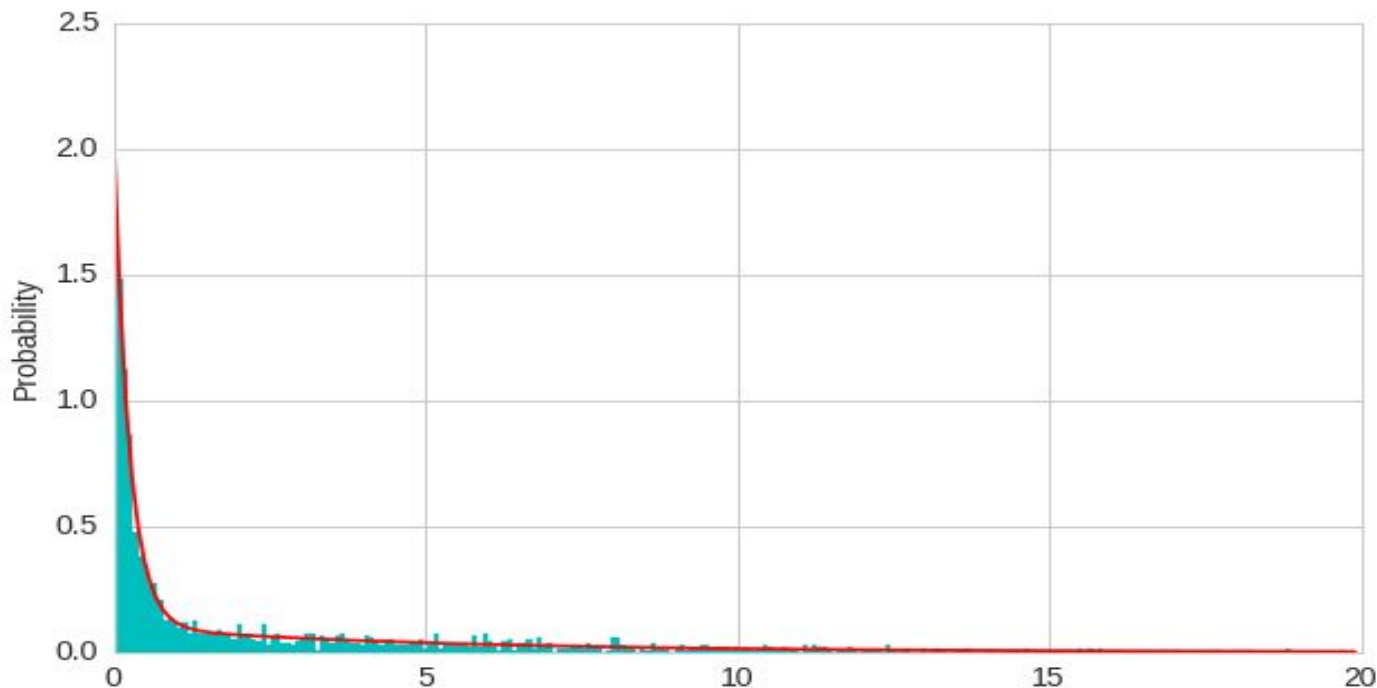


A single exponential distribution does not model this data well



```
model = ExponentialDistribution.from_samples(X)
```

A mixture of two exponentials models the data much better



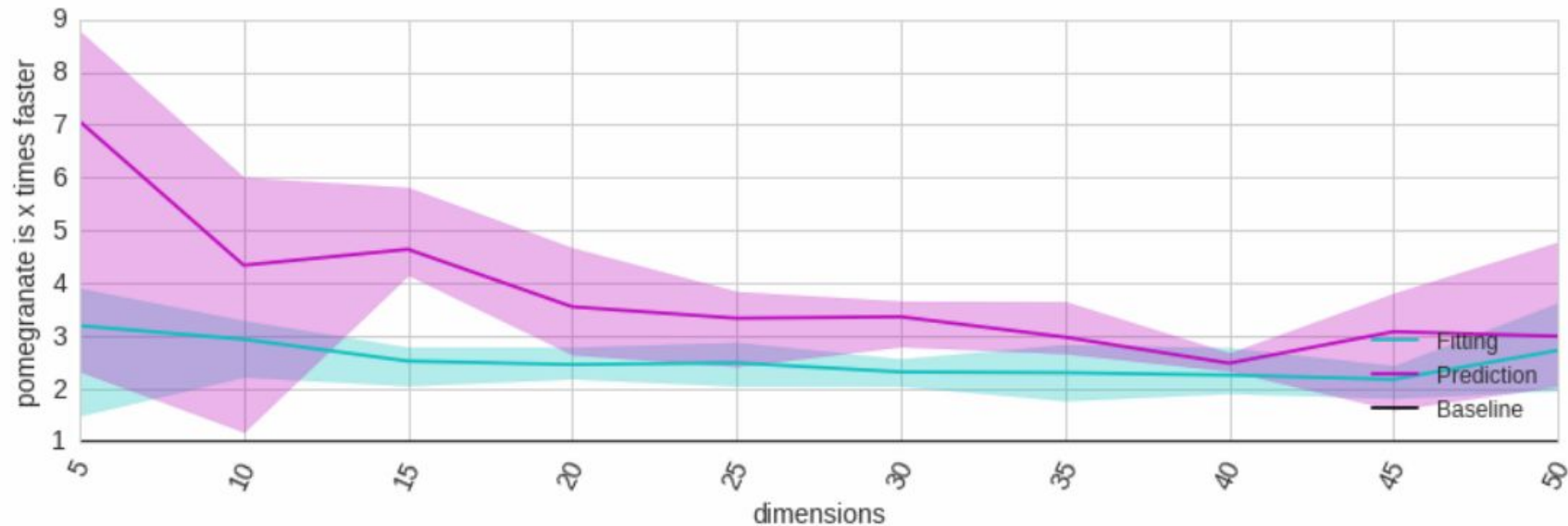
```
model = GeneralMixtureModel.from_samples(ExponentialDistribution, 2, X)
```

Heterogeneous mixtures are natively supported



```
model = GeneralMixtureModel.from_samples([ExponentialDistribution, UniformDistribution], 2, X)
```

general mixture models are faster than sklearn



Overview

The API

Major Models/Model Stacks

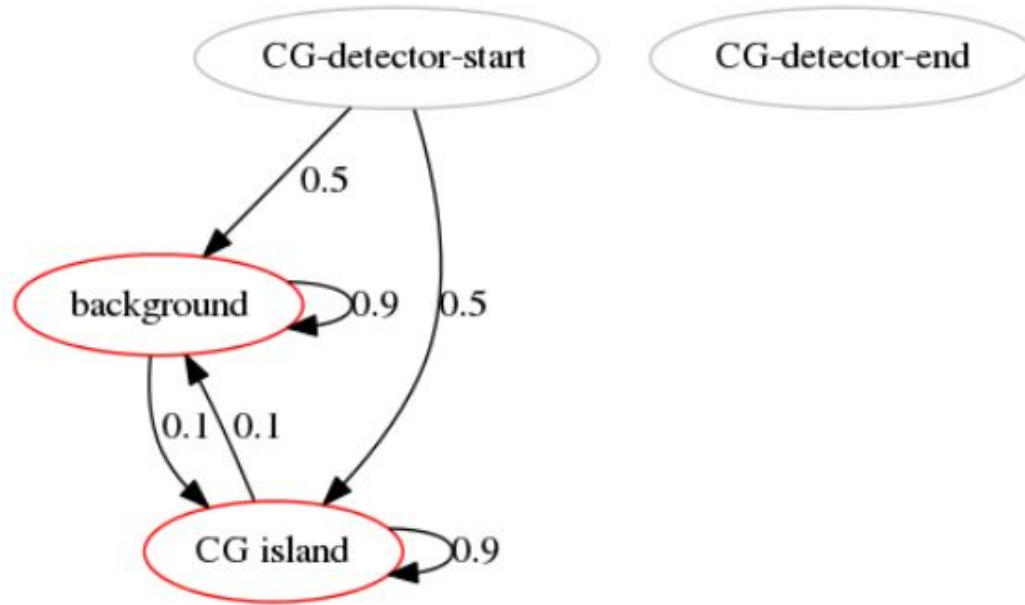
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CG enrichment detection HMM

GACTACGACTCGCGCTCGCGCGACGCGCTCGACATCATCGACACGACACTC



example: CG enrichment detector

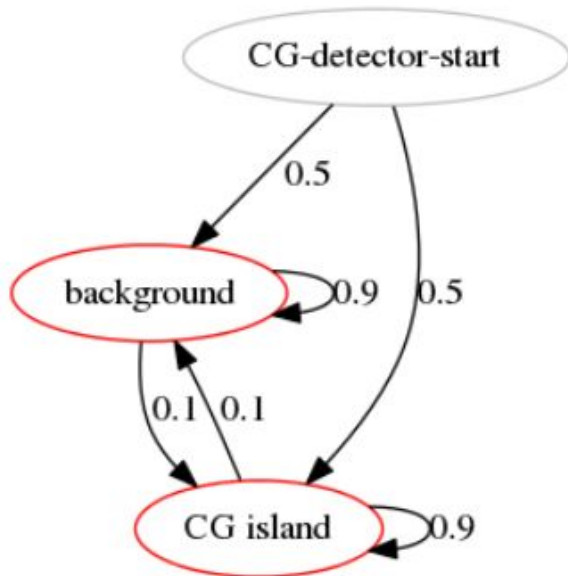
GACTACGACTCGCGCTCGCGCGACGCGCTCGACATCATCGACACGACACTC

```
d1 = DiscreteDistribution({'A': 0.25, 'C': 0.25, 'G': 0.25, 'T': 0.25})
d2 = DiscreteDistribution({'A': 0.10, 'C': 0.40, 'G': 0.40, 'T': 0.10})

s1 = State(d1, name="background")
s2 = State(d2, name="CG island")

hmm = HiddenMarkovModel("CG-detector")
hmm.add_states(s1, s2)
hmm.add_transition(hmm.start, s1, 0.5)
hmm.add_transition(hmm.start, s2, 0.5)
hmm.add_transition(s1, s1, 0.9)
hmm.add_transition(s1, s2, 0.1)
hmm.add_transition(s2, s1, 0.1)
hmm.add_transition(s2, s2, 0.9)
hmm.bake()
```

example: CG enrichment detector



```
d1 = DiscreteDistribution({'A': 0.25, 'C': 0.25, 'G': 0.25, 'T': 0.25})
d2 = DiscreteDistribution({'A': 0.10, 'C': 0.40, 'G': 0.40, 'T': 0.10})

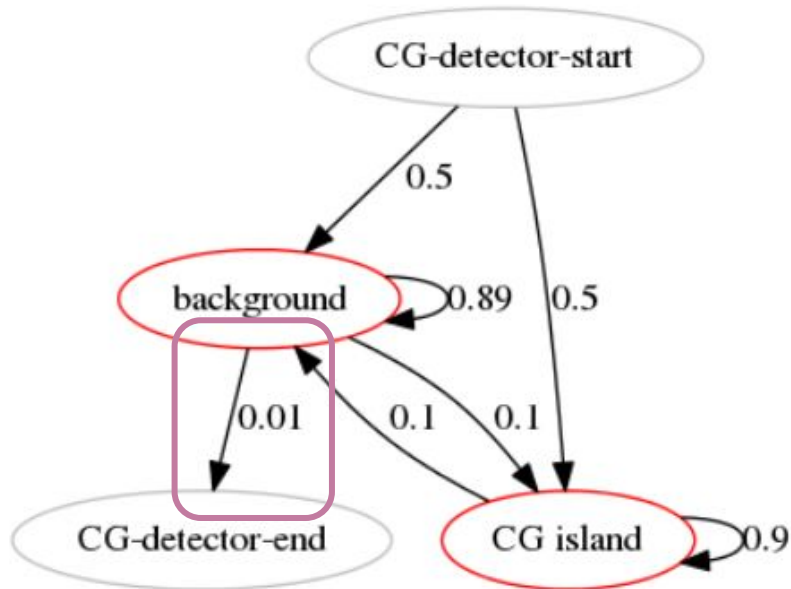
s1 = State(d1, name="background")
s2 = State(d2, name="CG island")

hmm = HiddenMarkovModel("CG-detector")
hmm.add_states(s1, s2)
hmm.add_transition(hmm.start, s1, 0.5)
hmm.add_transition(hmm.start, s2, 0.5)
hmm.add_transition(s1, s1, 0.9)
hmm.add_transition(s1, s2, 0.1)
hmm.add_transition(s2, s1, 0.1)
hmm.add_transition(s2, s2, 0.9)
hmm.bake()
```

```
sequence: CGACTACTGACTACTCGCGCGCACGCGGCACGCGTGCCGTCTATACTGCGCATACGGC  
hmm pred: 1111111111111111000000000000000000000000000000000000000000000000
```

```
hmm state 0: CG island
hmm state 1: background
```


example: CG enrichment detector



```
hmm = HiddenMarkovModel("CG-detector")
hmm.add_states(s1, s2)
hmm.add_transition(hmm.start, s1, 0.5)
hmm.add_transition(hmm.start, s2, 0.5)
hmm.add_transition(s1, s1, 0.89)
hmm.add_transition(s1, s2, 0.10)
hmm.add_transition(s1, hmm.end, 0.01)
hmm.add_transition(s2, s1, 0.1)
hmm.add_transition(s2, s2, 0.9)
hmm.bake()
```

hidden markov models

Feature	pomegranate	hmmlearn
Graph Structure		
Silent States	✓	
Optional Explicit End State	✓	
Sparse Implementation	✓	
Arbitrary Emissions Allowed on States	✓	
Discrete/Gaussian/GMM Emissions	✓	✓
Large Library of Other Emissions	✓	
Build Model from Matrices	✓	✓
Build Model Node-by-Node	✓	
Serialize to JSON	✓	
Serialize using Pickle/Joblib	✓	✓

Algorithms		
Priors		✓
Sampling	✓	✓
Log Probability Scoring	✓	✓
Forward-Backward Emissions	✓	✓
Forward-Backward Transitions	✓	
Viterbi Decoding	✓	✓
MAP Decoding	✓	✓
Baum-Welch Training	✓	✓
Viterbi Training	✓	
Labeled Training	✓	
Tied Emissions	✓	
Tied Transitions	✓	
Emission Inertia	✓	
Transition Inertia	✓	
Emission Freezing	✓	✓
Transition Freezing	✓	✓
Multi-threaded Training	✓	

hidden markov models

Feature	pomegranate	hmmlearn
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MAP Decoding	✓	✓
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Viterbi Training	✓	
Labeled Training	✓	
Tied Emissions	✓	
Tied Transitions	✓	
Emission Inertia	✓	
Transition Inertia	✓	
Emission Freezing	✓	✓
Transition Freezing	✓	✓
Multi-threaded Training	✓	

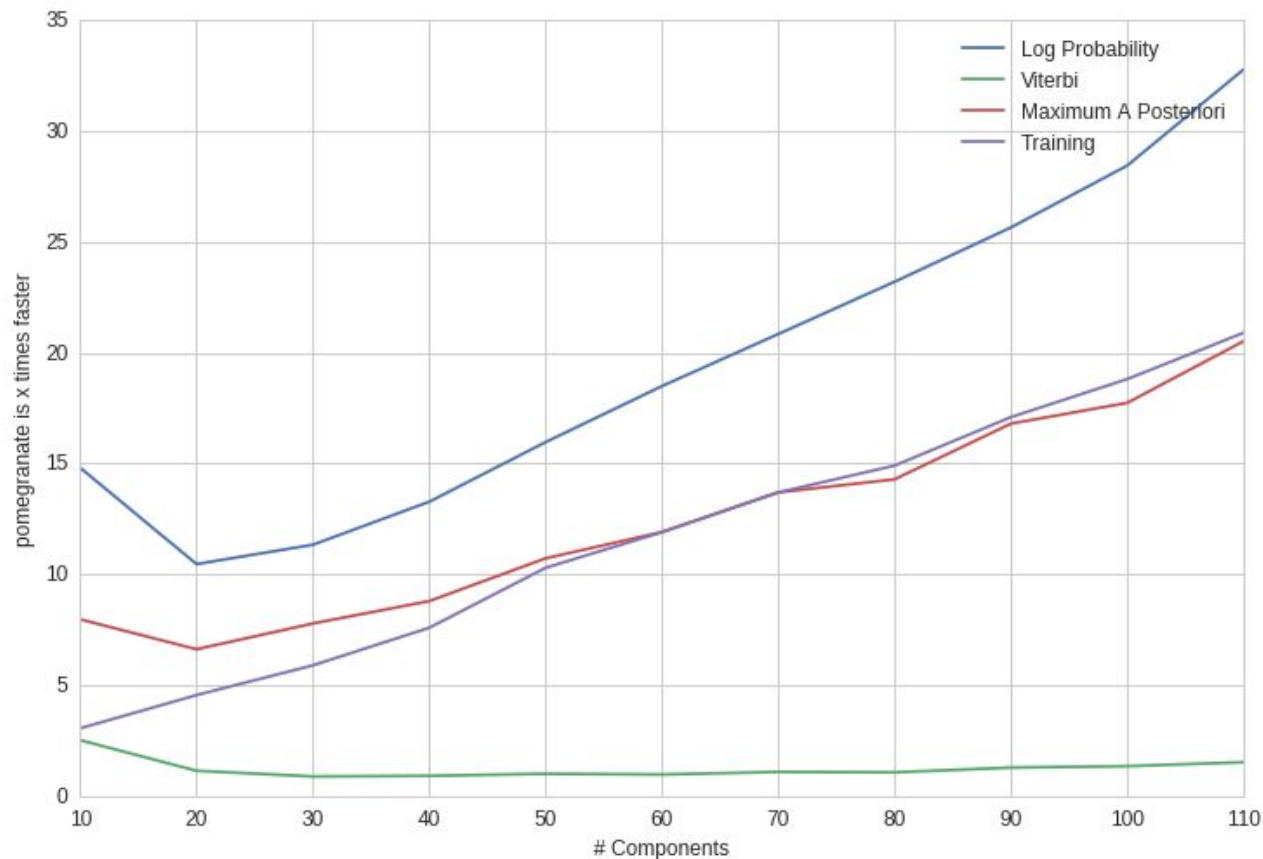
example: GMM-HMM

```
d1 = GeneralMixtureModel([NormalDistribution(5, 2), NormalDistribution(5, 4)])
d2 = GeneralMixtureModel([NormalDistribution(15, 1), NormalDistribution(15, 5)])

s1 = State(d1, name="GMM1")
s2 = State(d2, name="GMM2")

model = HiddenMarkovModel()
model.add_states(s1, s2)
model.add_transition(model.start, s1, 0.75)
model.add_transition(model.start, s2, 0.25)
model.add_transition(s1, s1, 0.85)
model.add_transition(s1, s2, 0.15)
model.add_transition(s2, s2, 0.90)
model.add_transition(s2, s1, 0.10)
model.bake()
```

hidden markov models are faster than hmmlearn



Overview

The API

Major Models/Model Stacks

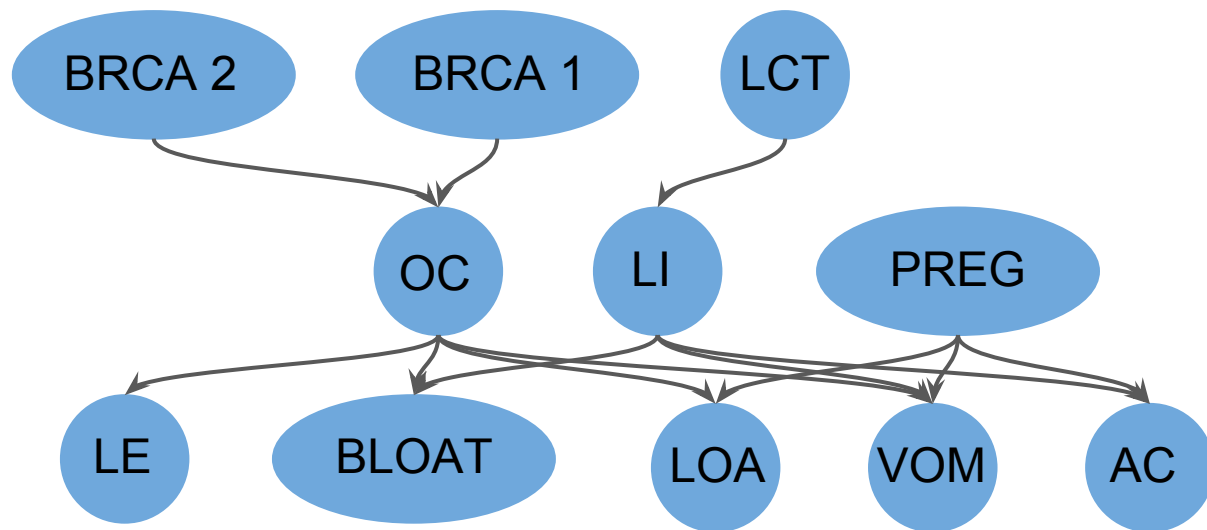
1. General Mixture Models
2. Hidden Markov Models
3. **Bayesian Networks**
4. Bayes Classifiers

Parallelization

Finale: Train a mixture of hidden markov models in parallel

Bayesian networks

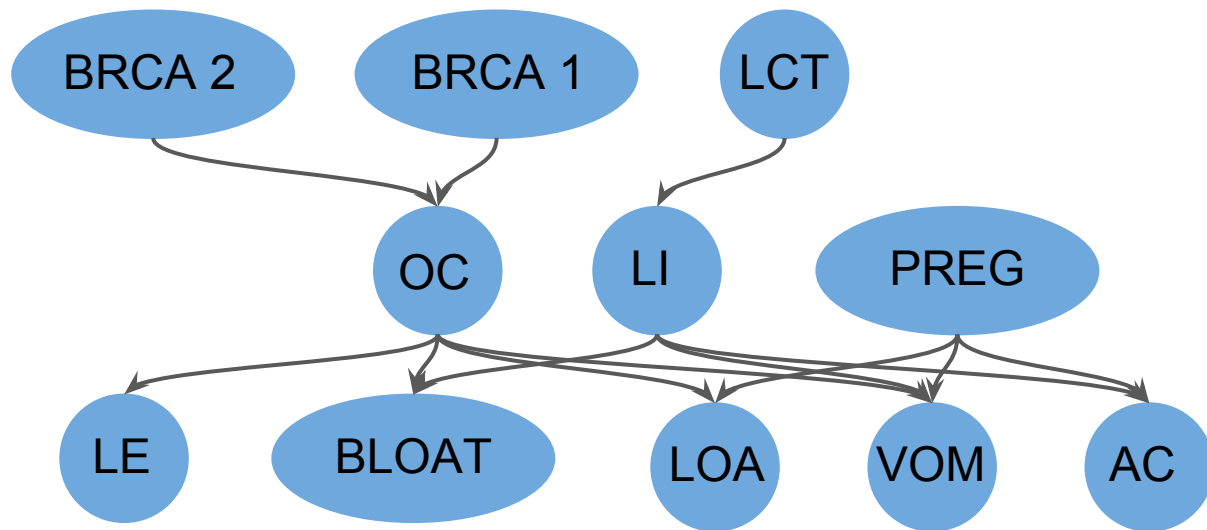
Bayesian networks are powerful inference tools which define a dependency structure between variables.



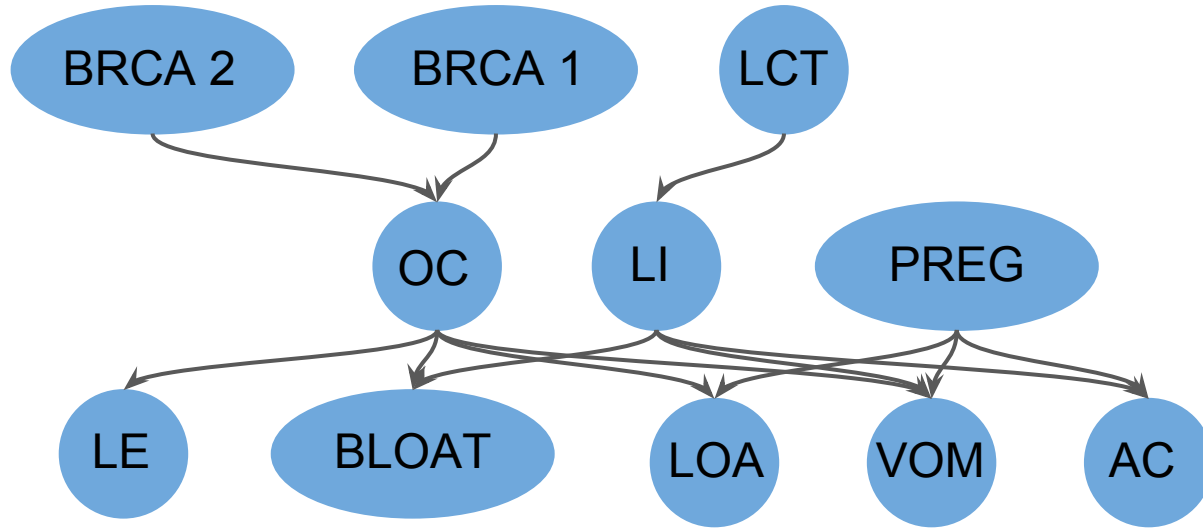
Bayesian networks

Two main non-trivial tasks:

- (1) Inference given incomplete information
- (2) Learning the dependency structure from data



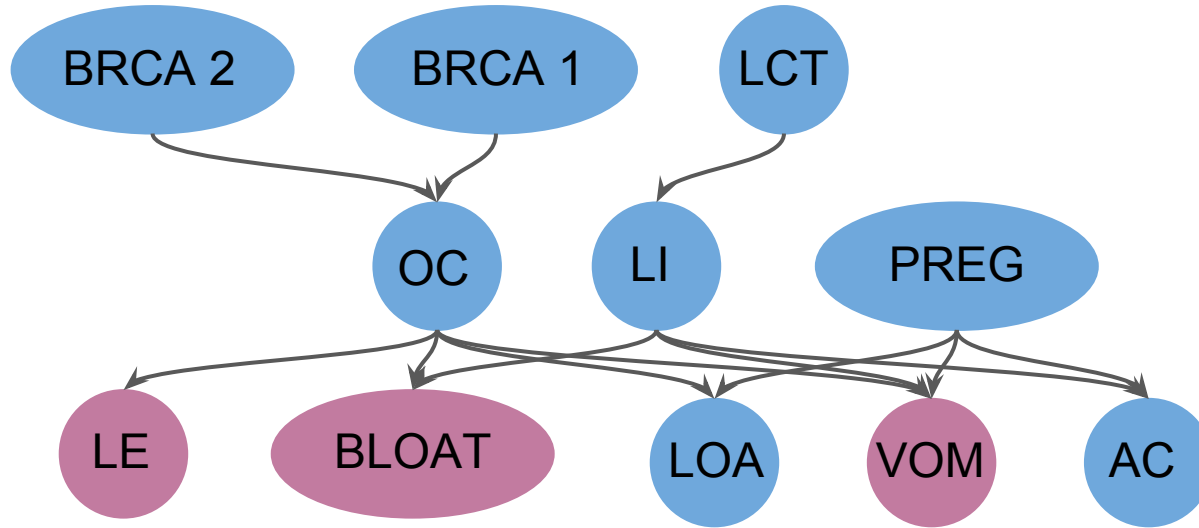
Inference given incomplete information



```
d = model.predict_proba()
print "\t".join( "{:7}".format(state.name) for state in model.states )
print "\t".join( "{:4.2}".format(model.parameters[0][1]) for model in d )
```

BRCA1	BRCA2	LCT	OC	LI	PREG	LE	BLOAT	LOA	VOM	AC
0.001	0.015	0.05	0.005	0.05	0.1	0.014	0.16	0.017	0.091	0.15

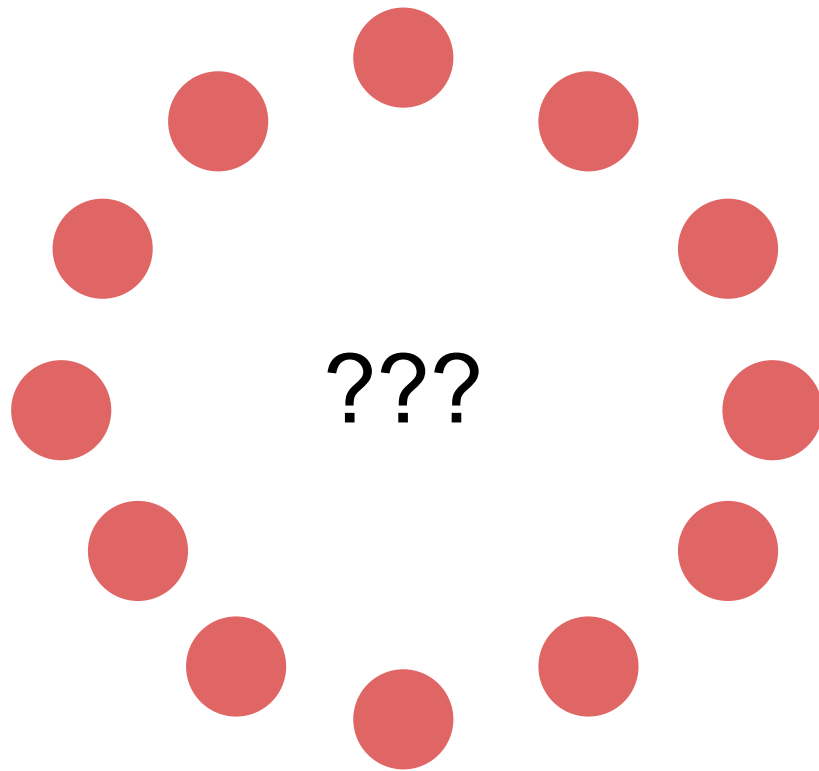
Inference given incomplete information



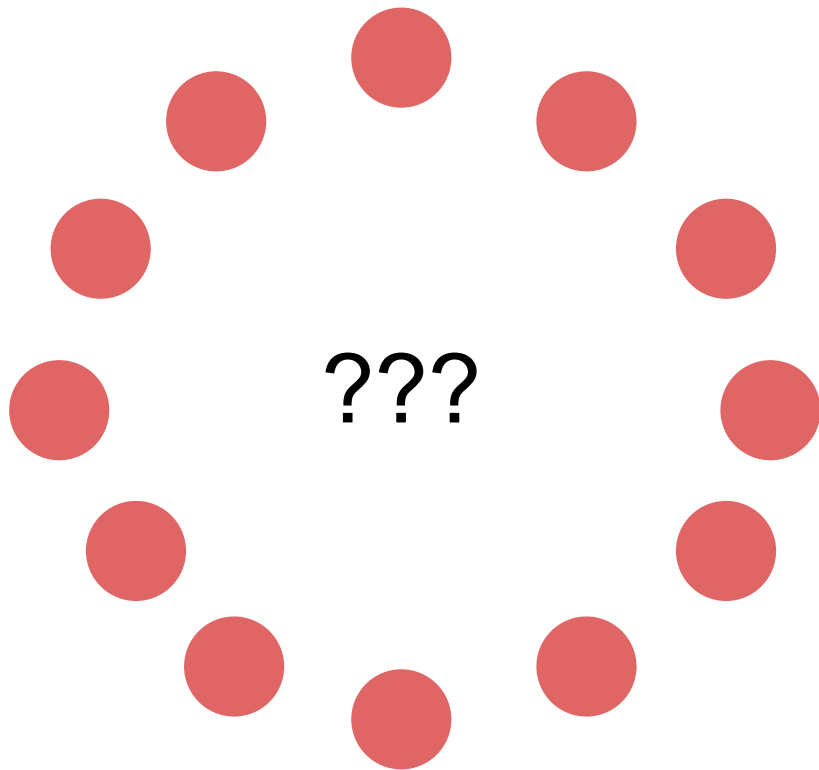
```
d = model.predict_proba({'VOM' : 1, 'BLOAT' : 1, 'LE' : 1})
print "\t".join( "{:7}".format(state.name) for state in model.states )
print "\t".join( "{:4.2}".format(model.parameters[0][1]) for model in d )
```

BRCA1	BRCA2	LCT	OC	LI	PREG	LE	BLOAT	LOA	VOM	AC
0.056	0.68	0.087	0.91	0.096	0.2	1.0	1.0	0.52	1.0	0.24

Sometimes we want to learn structure from data



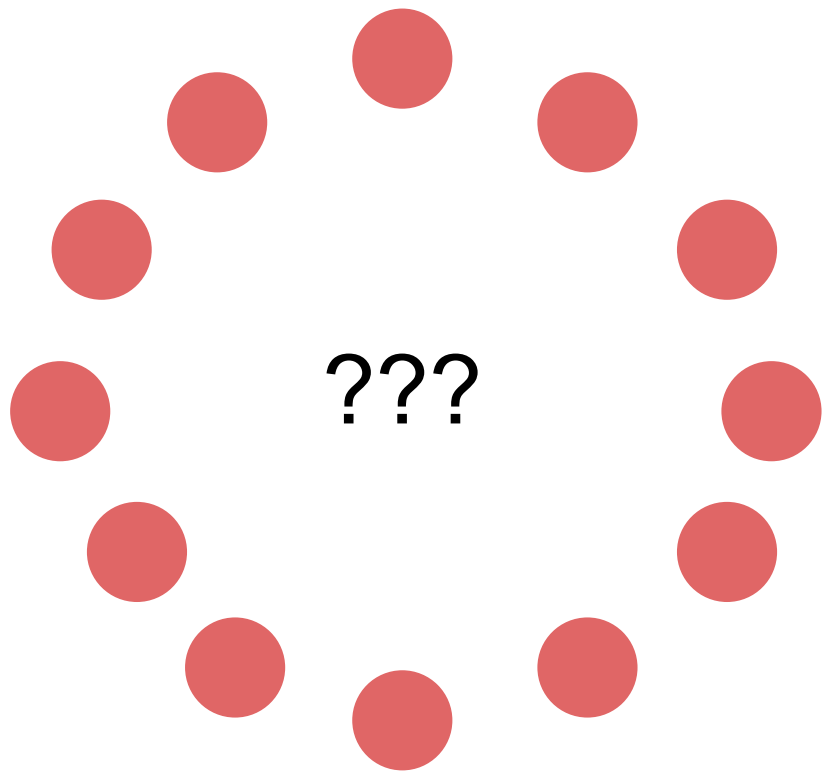
Sometimes we want to learn structure from data



Three primary ways:

- “Search and score” / Exact
- “Constraint Learning” / PC
- Heuristics

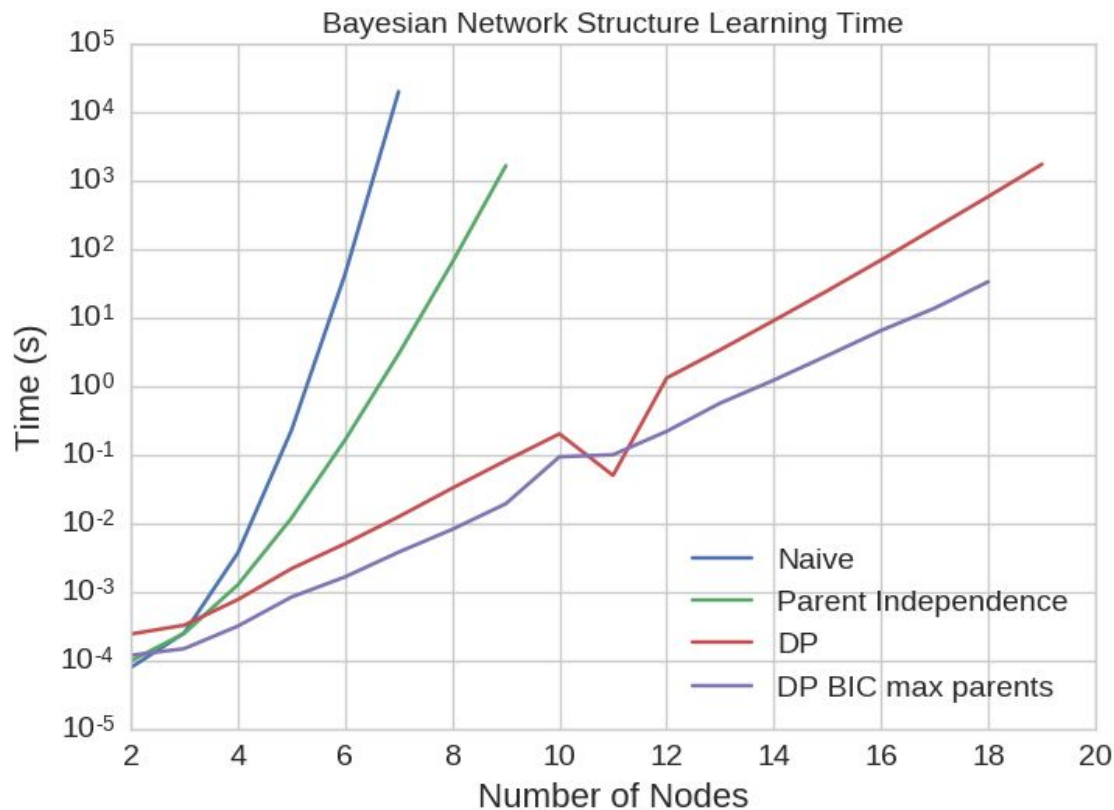
Sometimes we want to learn structure from data



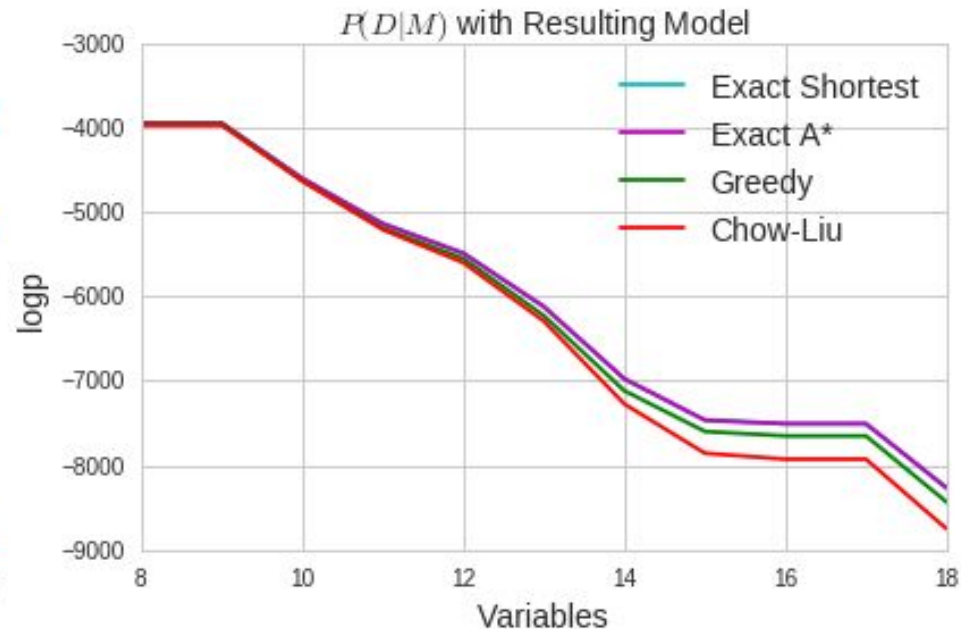
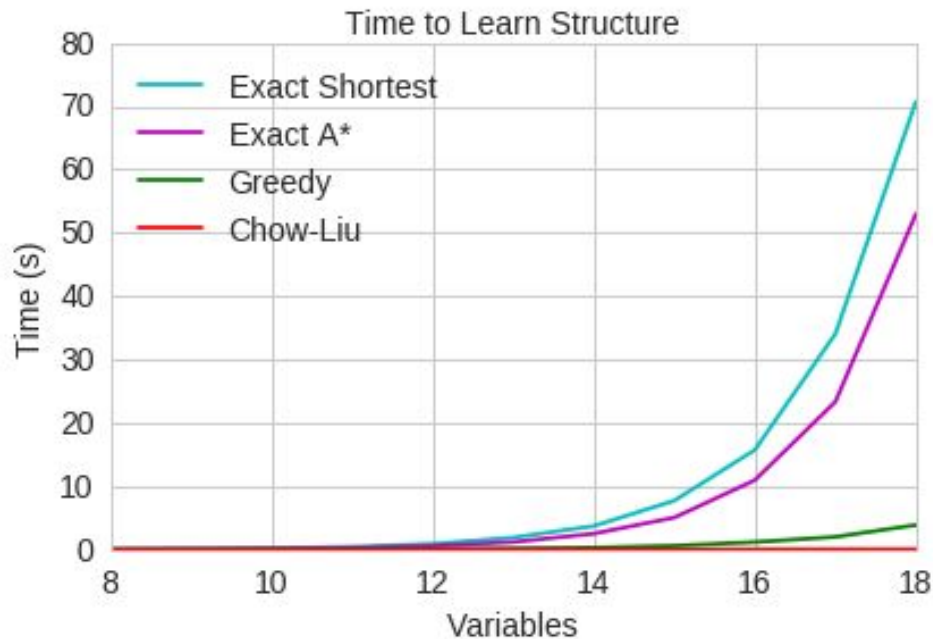
pomegranate supports

- “Search and score” / Exact
- “Constraint Learning” / PC
- Heuristics

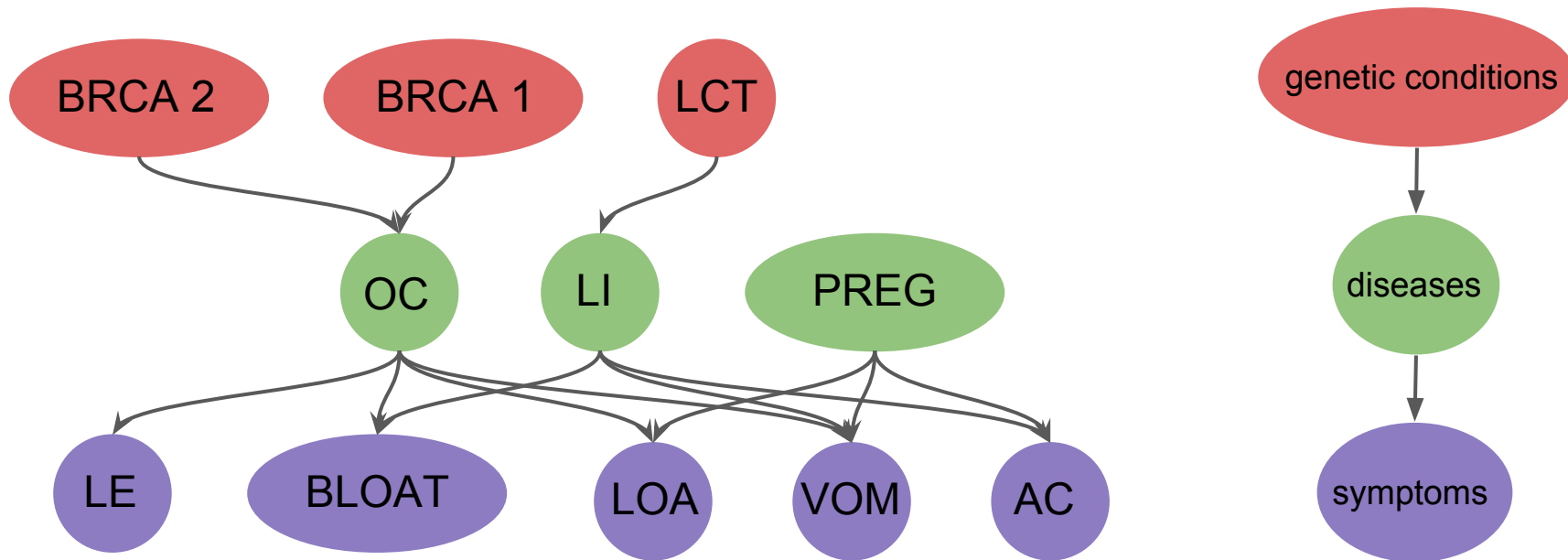
Structure learning can be super-exponential in time



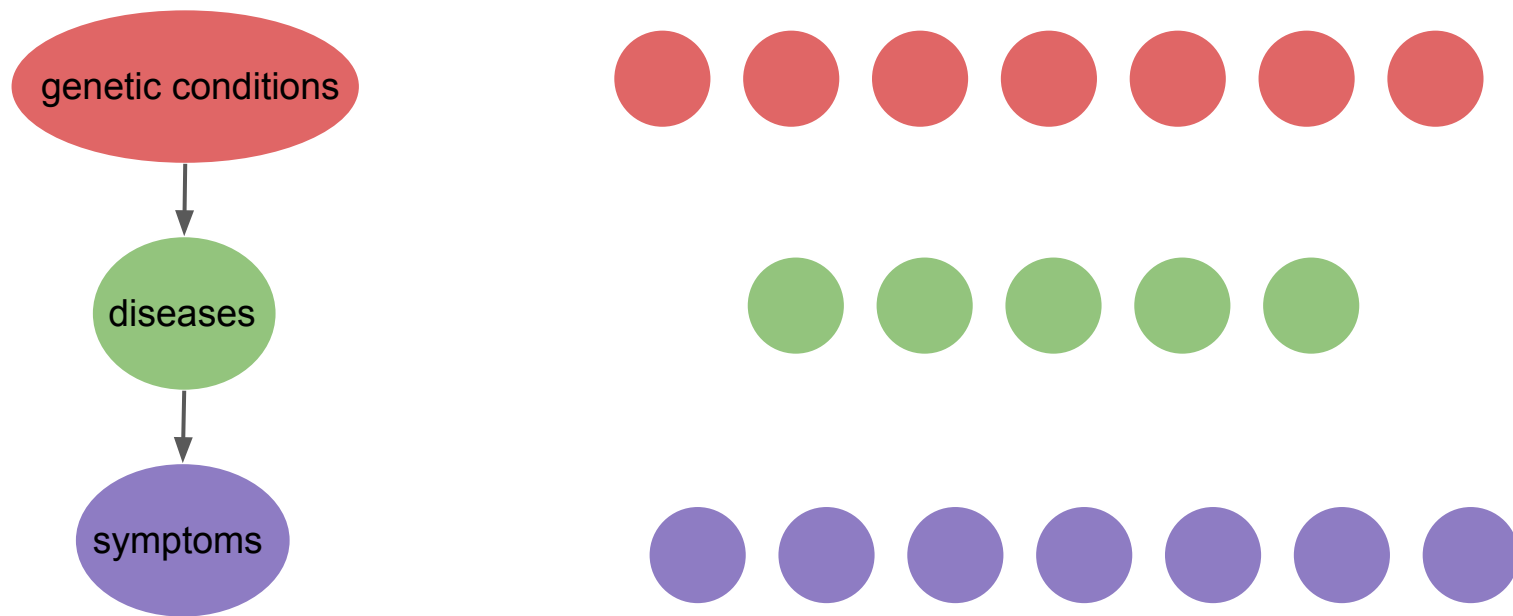
pomegranate supports 4 structure learning algos



Constraint graphs can merge expert knowledge with data

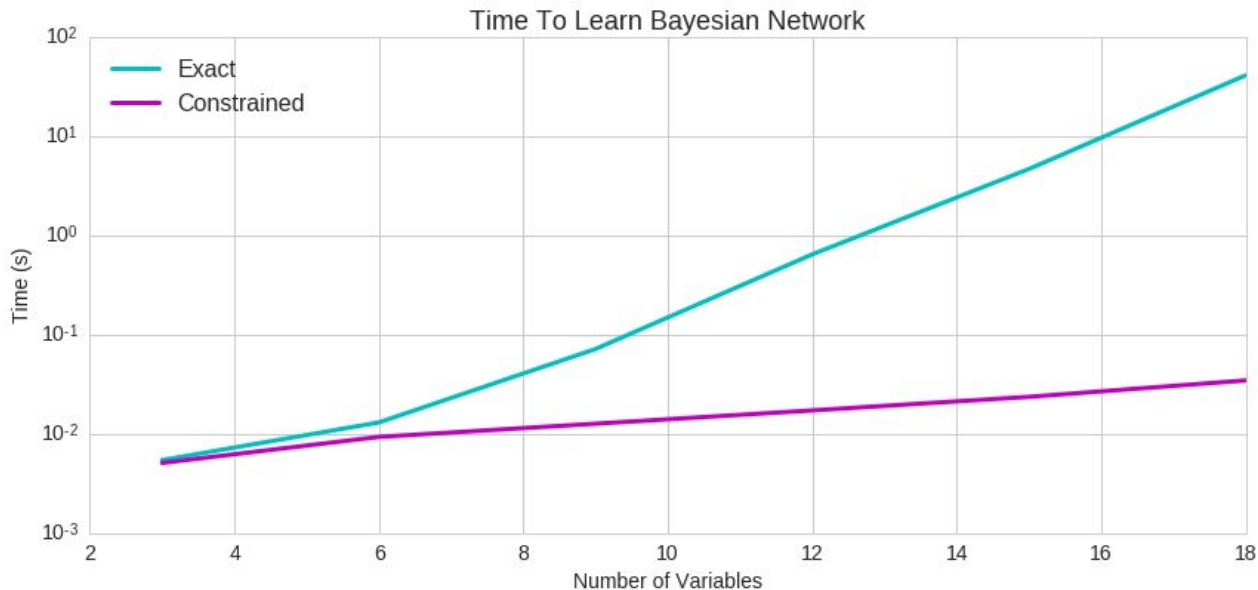
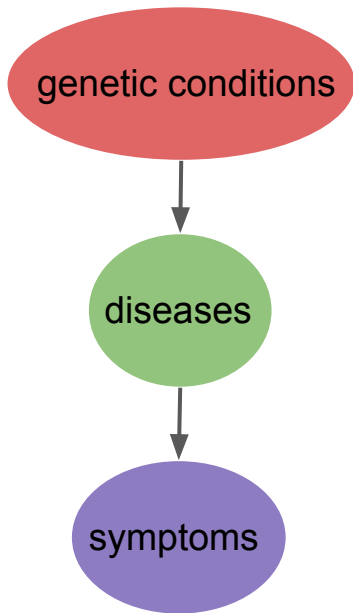


Structure learning with Constraint Graphs

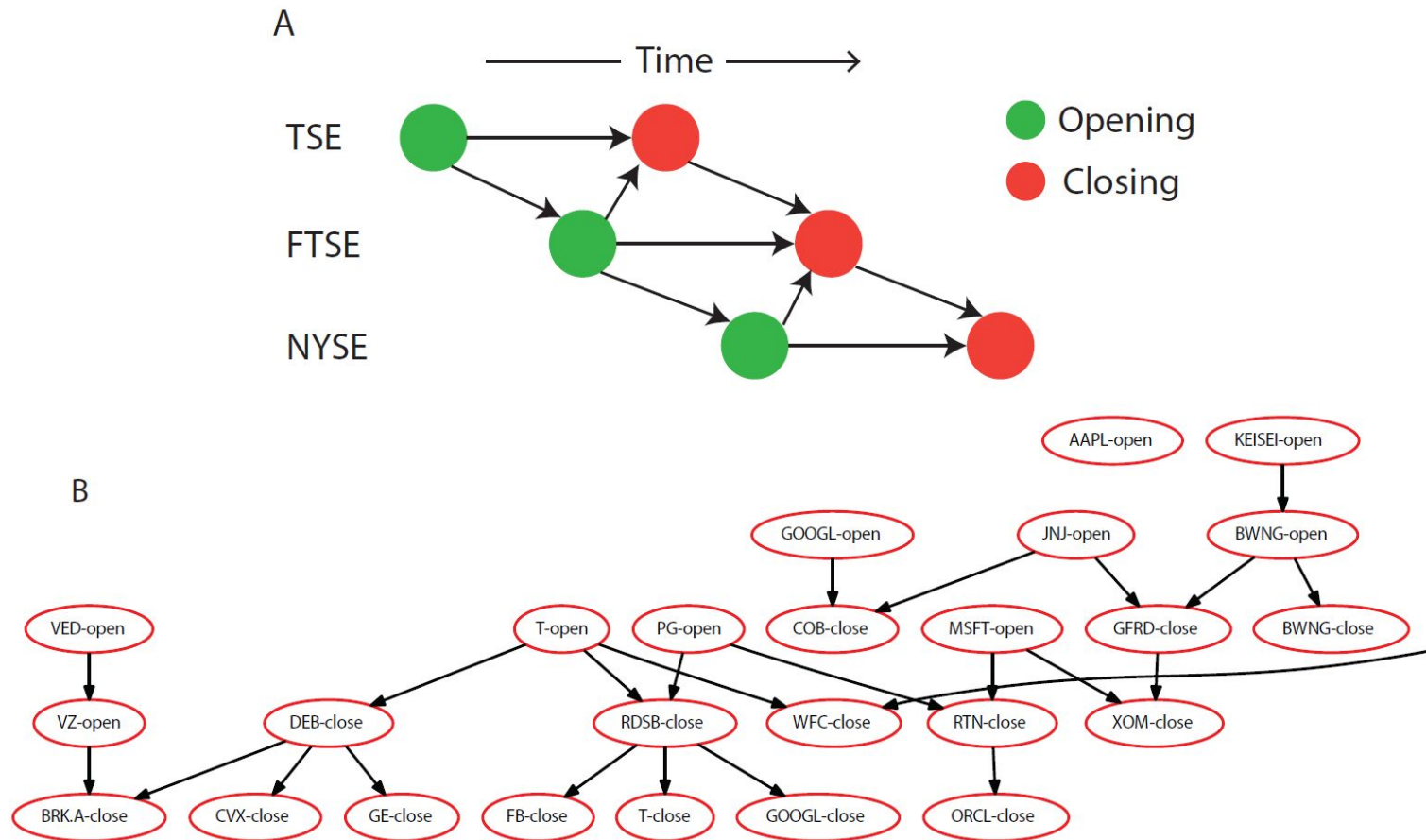


Structure learning with constraint graphs

Constraint graphs can also encode possible dependencies as layers.



Constraint graphs can model the global stock market



Overview

The API

Major Models/Model Stacks

1. General Mixture Models
2. Hidden Markov Models
3. Bayesian Networks
4. Bayes Classifiers

Parallelization

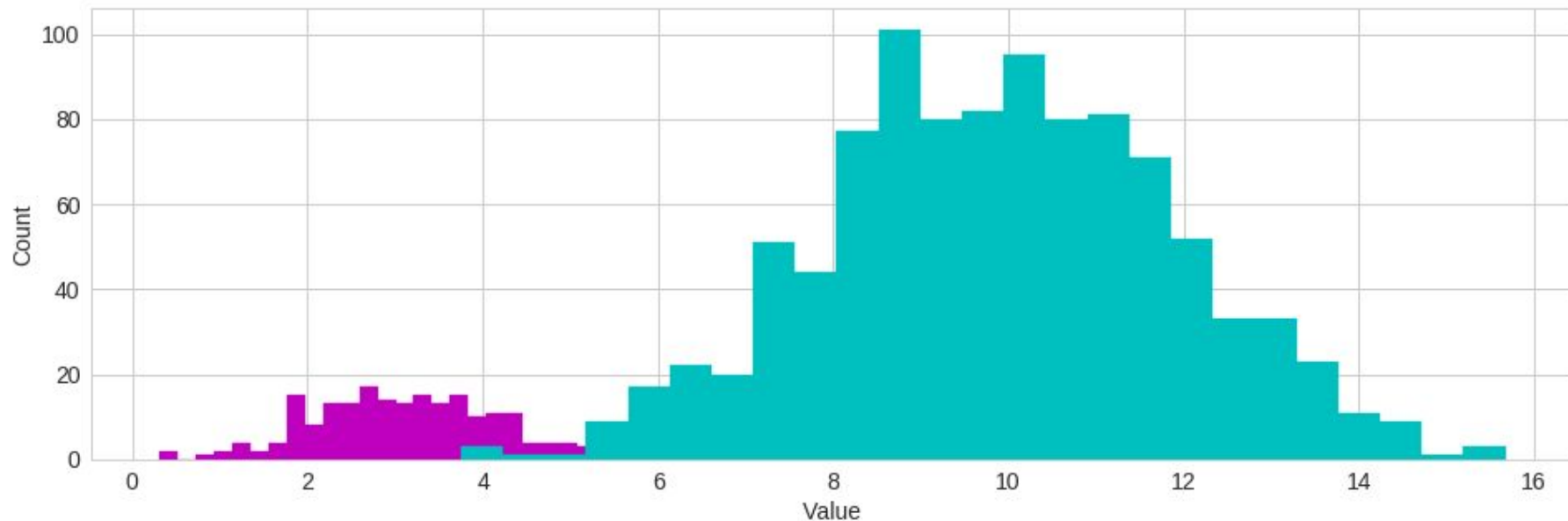
Finale: Train a mixture of hidden markov models in parallel

Bayes' Rule

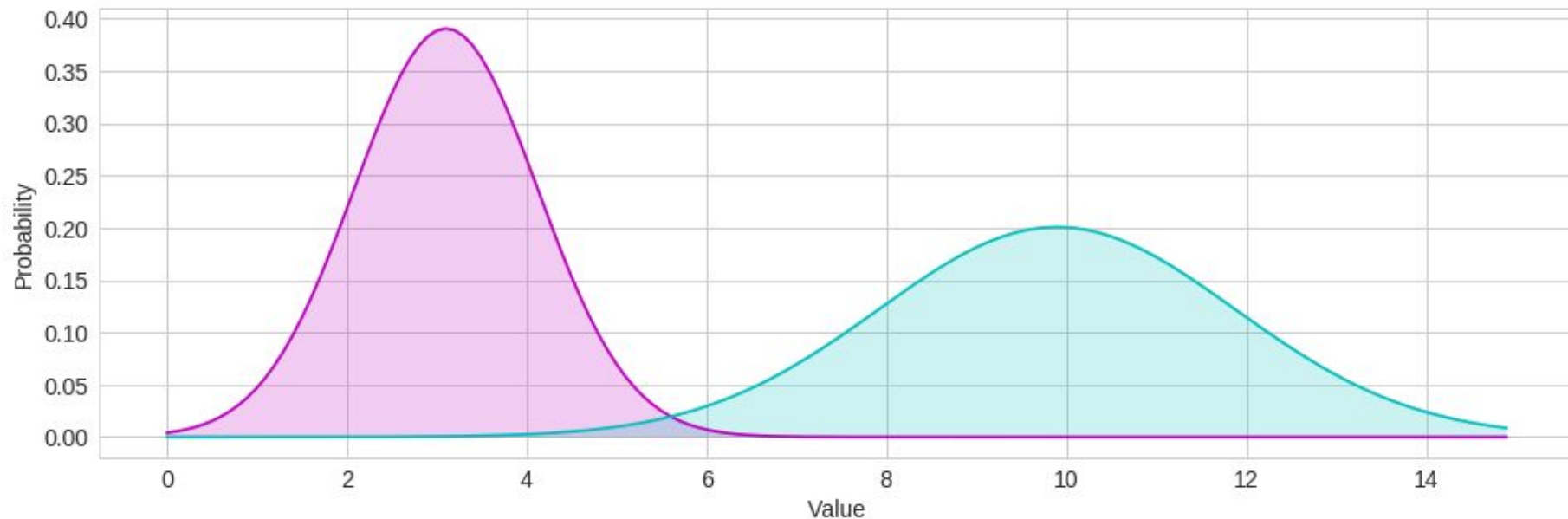
$$P(M|D) = P(D|M)P(M) / P(D)$$

$$\text{Posterior} = \text{Likelihood} * \text{Prior} / \text{Normalization}$$

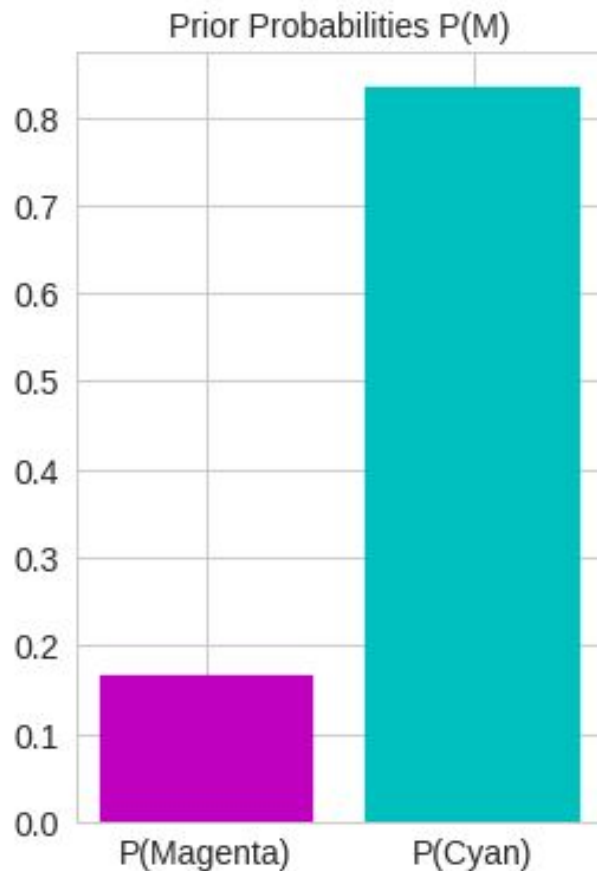
Let's build a simple classifier on this data



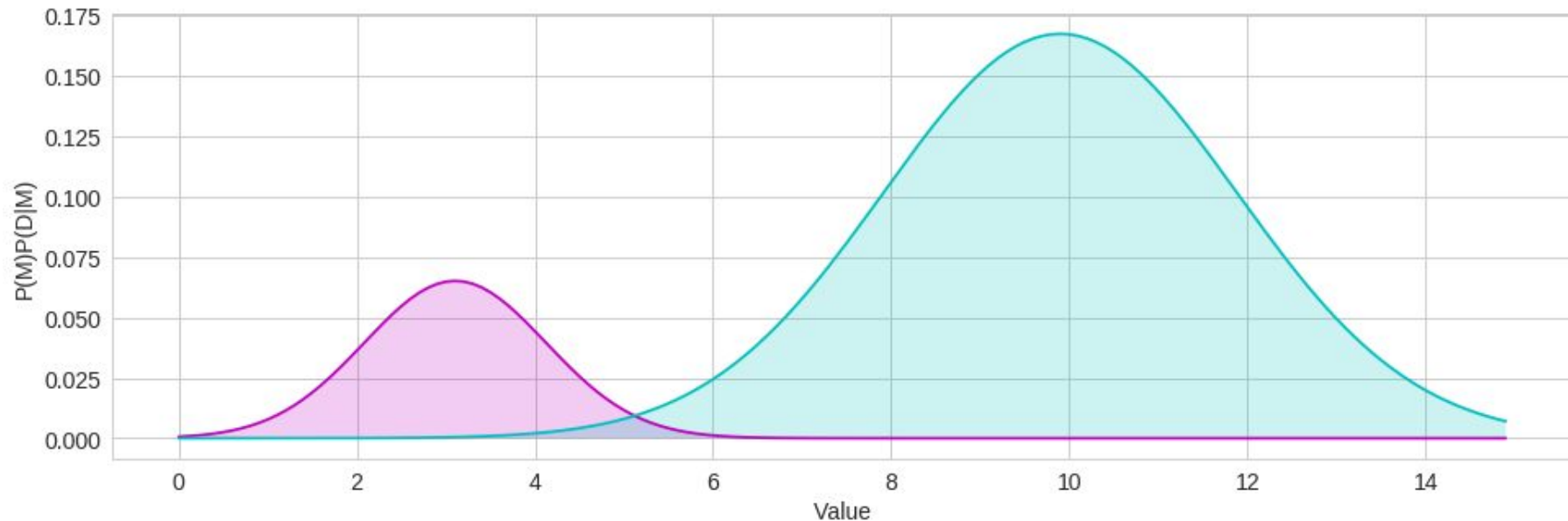
The likelihood function itself ignores class imbalance



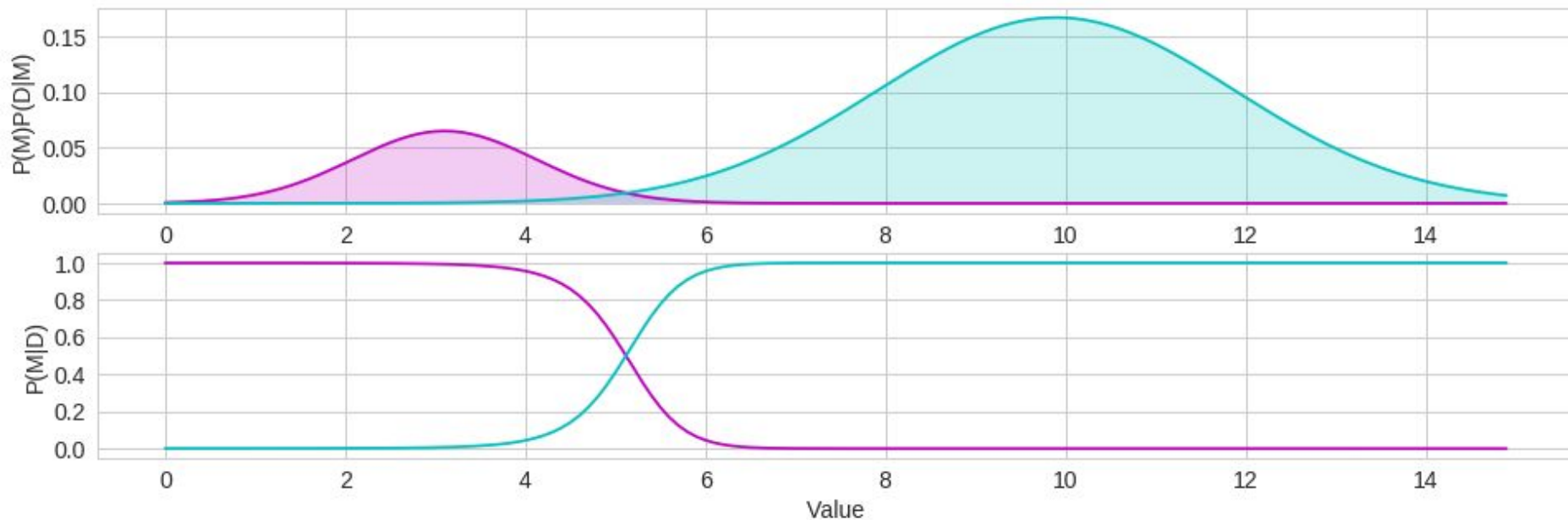
The prior probabilities can model class imbalance



The posterior models the original data more faithfully



The ratio of the posterior is a good classifier



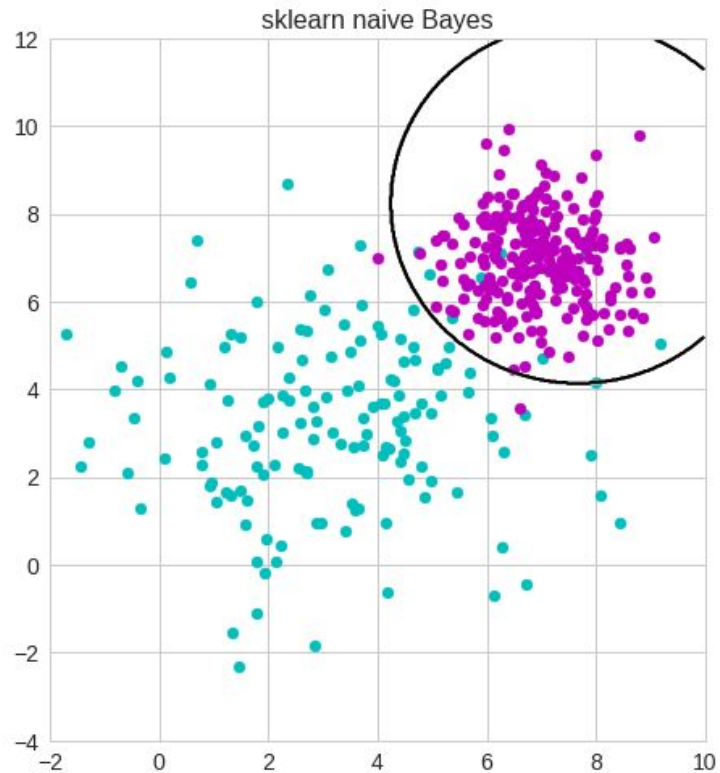
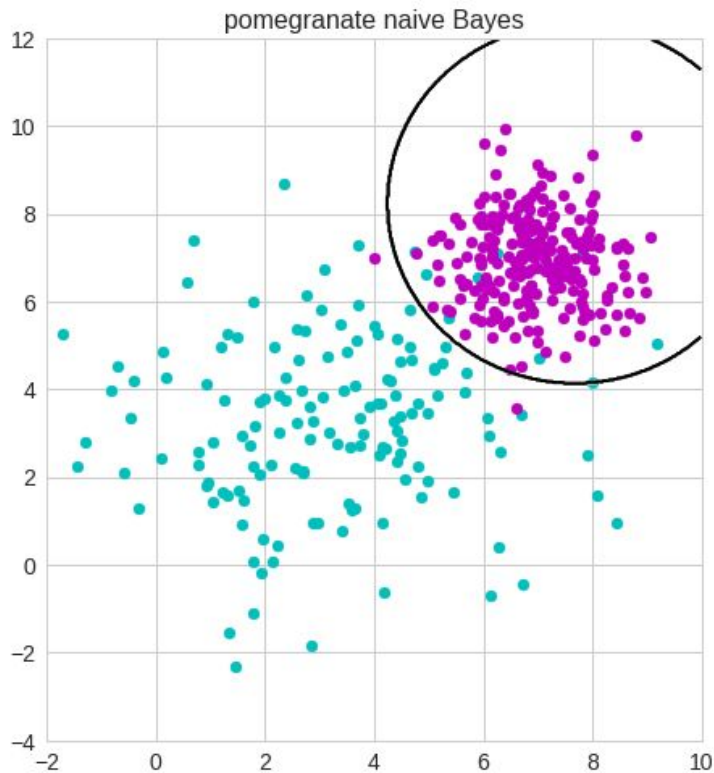
```
model = NaiveBayes.from_samples(NormalDistribution, X, y)
posteriors = model.predict_proba(idxs)
```

Naive Bayes assumes all dimensions are independent

$$P(M|D) = \prod P(D|M) P(M) / P(D)$$

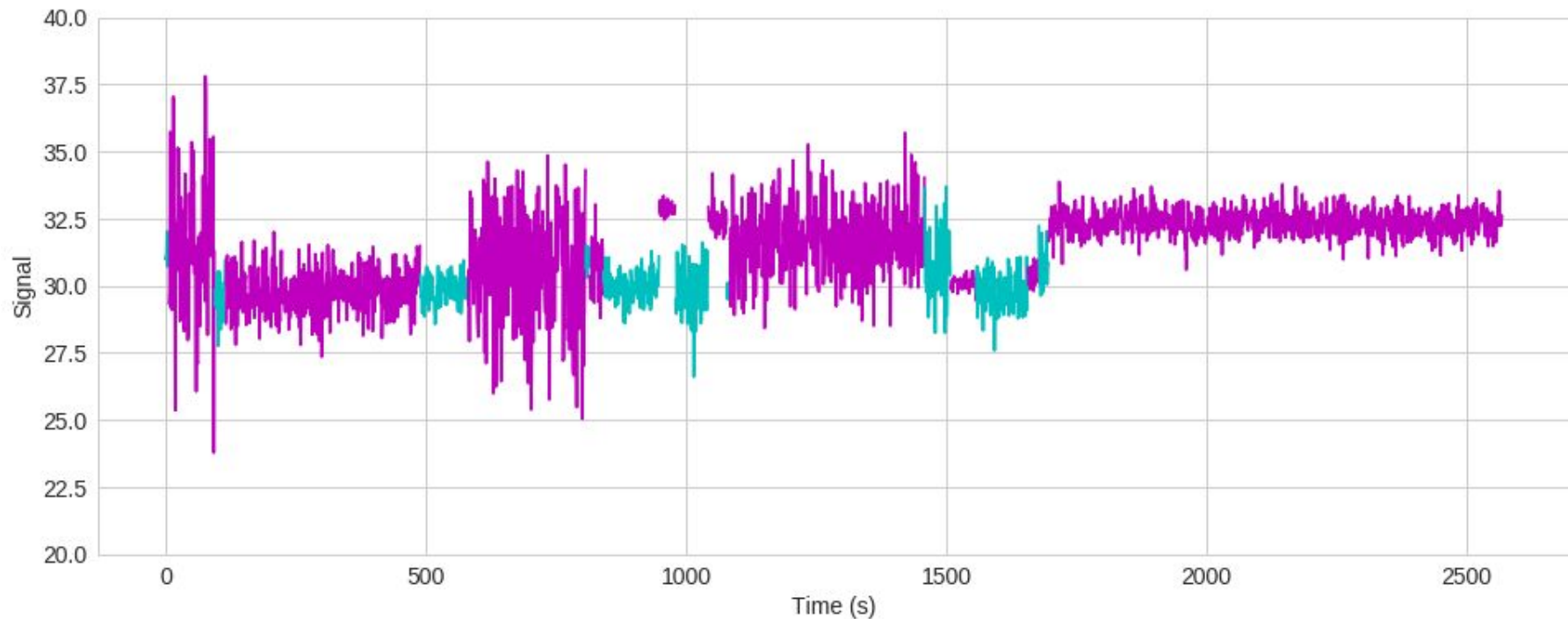
$$\text{Posterior} = \text{Likelihood} * \text{Prior} / \text{Normalization}$$

Gaussian naive Bayes produces spherical distributions

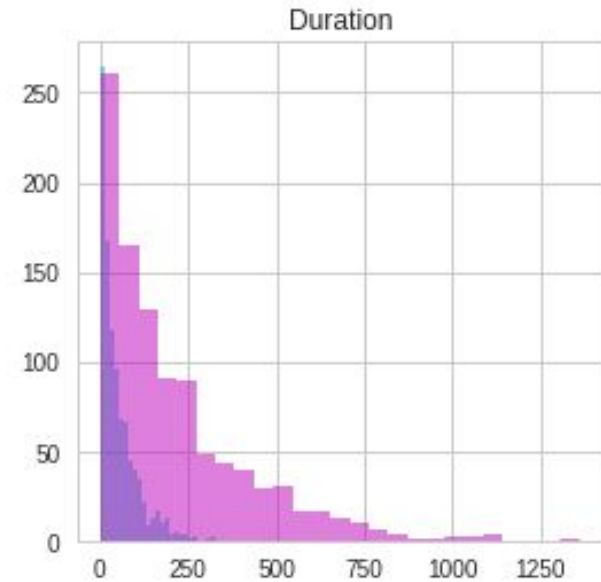
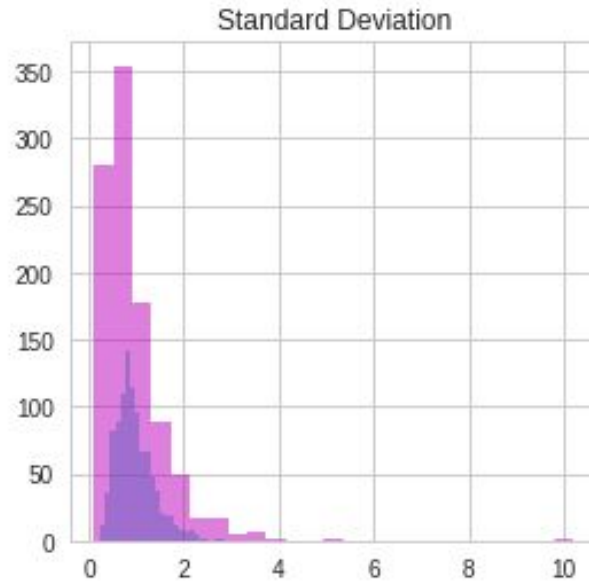
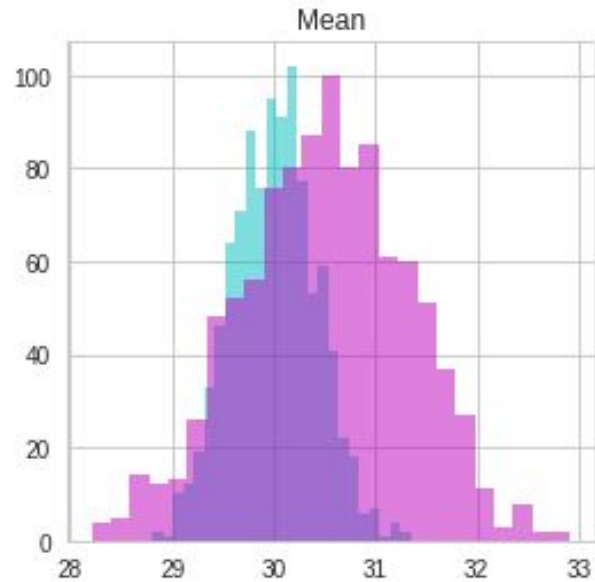


`model = NaiveBayes.from_samples(NormalDistribution, X, y)`

Naive Bayes does not need to be homogenous



Different features fall under different distributions



Explicitly modeling these distributions yields better classifiers

```
model = NaiveBayes.from_samples(NormalDistribution, X_train, y_train)
print "Gaussian Naive Bayes: ", (model.predict(X_test) == y_test).mean()
```

```
clf = GaussianNB().fit(X_train, y_train)
print "sklearn Gaussian Naive Bayes: ", (clf.predict(X_test) == y_test).mean()
```

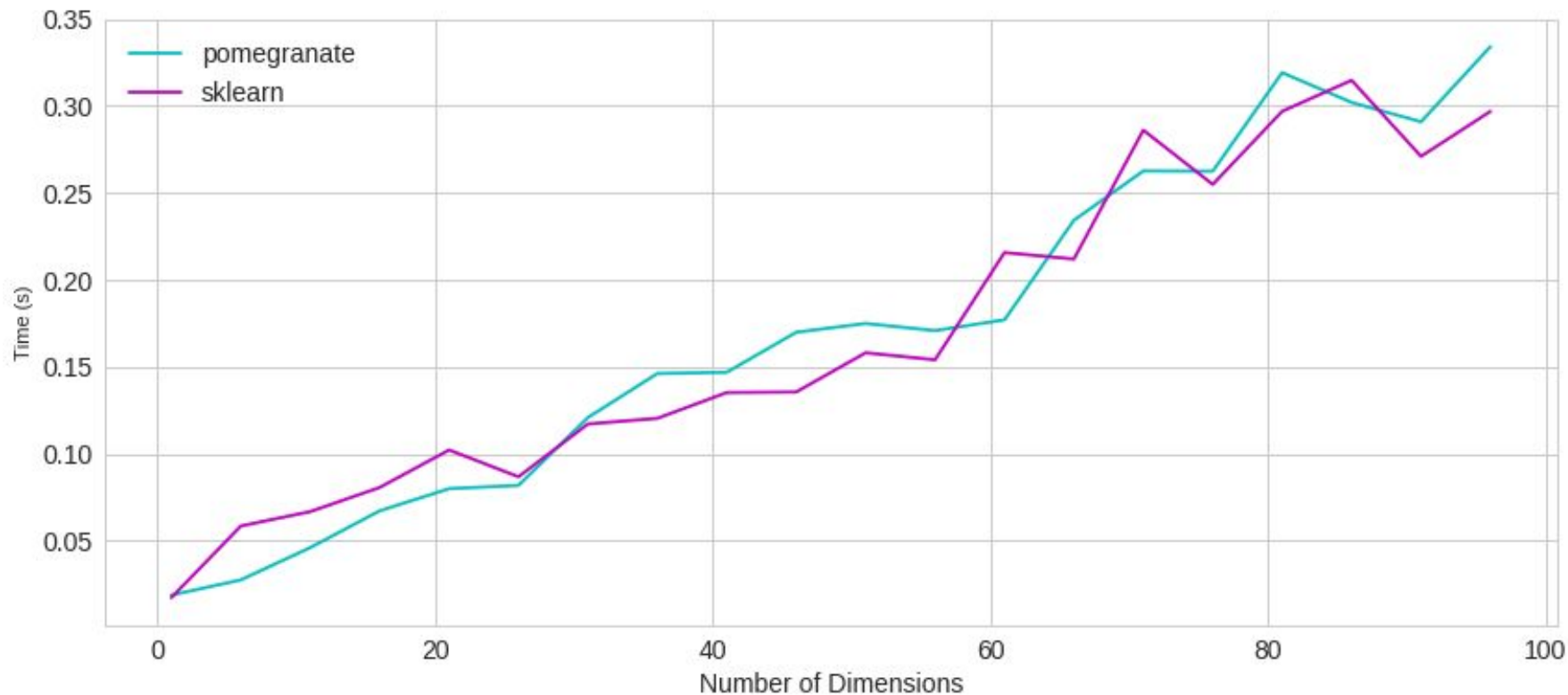
```
model = NaiveBayes.from_samples([NormalDistribution, LogNormalDistribution,
ExponentialDistribution], X_train, y_train)
print "Heterogeneous Naive Bayes: ", (model.predict(X_test) == y_test).mean()
```

Gaussian Naive Bayes: 0.798

sklearn Gaussian Naive Bayes: 0.798

Heterogeneous Naive Bayes: 0.844

pomegranate is just as fast as sklearn

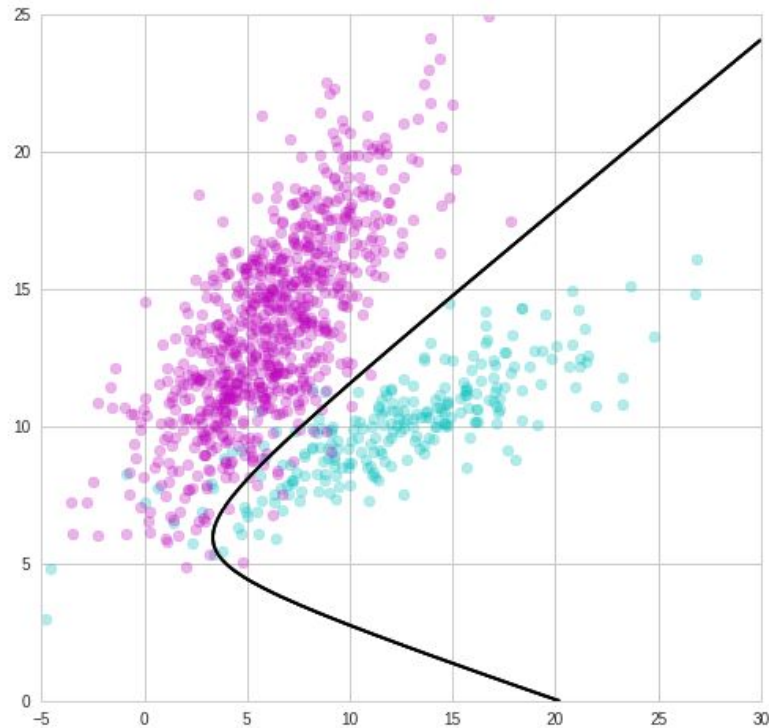
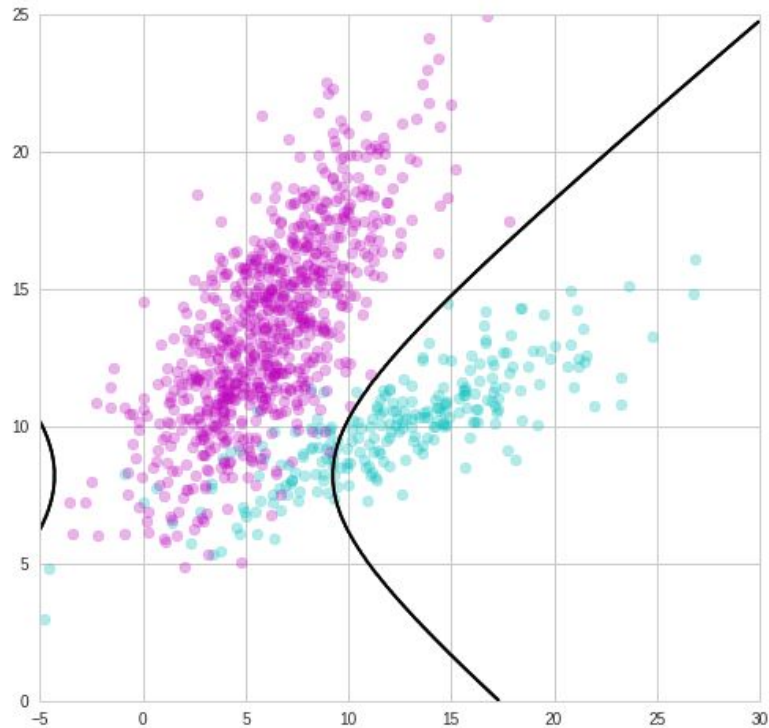


Bayes Classifiers are more general than naive Bayes

$$P(M|D) = P(D|M) P(M) / P(D)$$

$$\text{Posterior} = \text{Likelihood} * \text{Prior} / \text{Normalization}$$

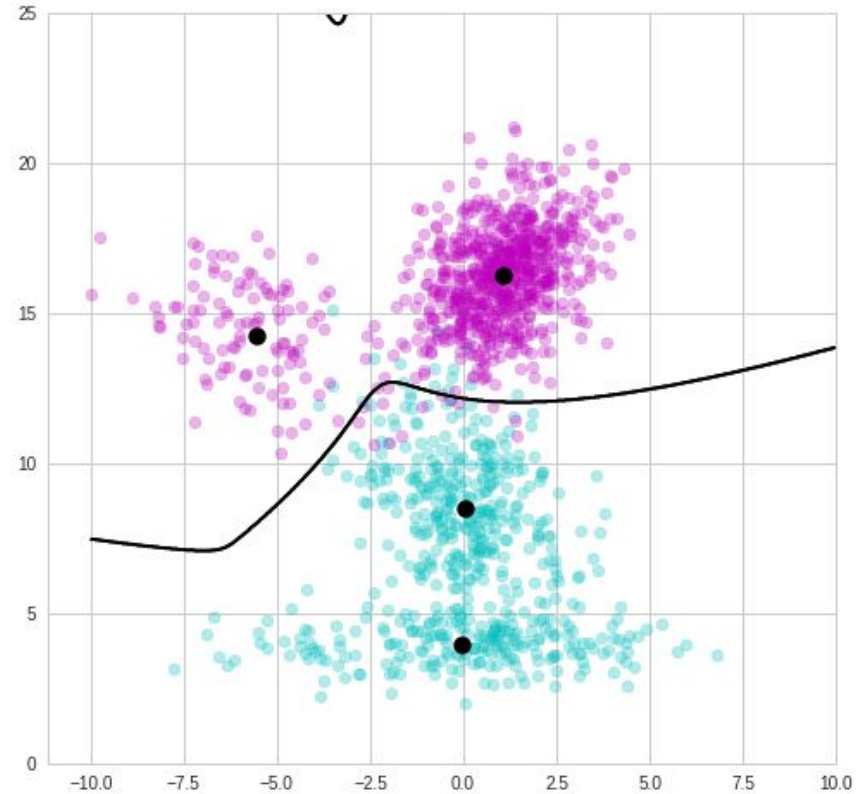
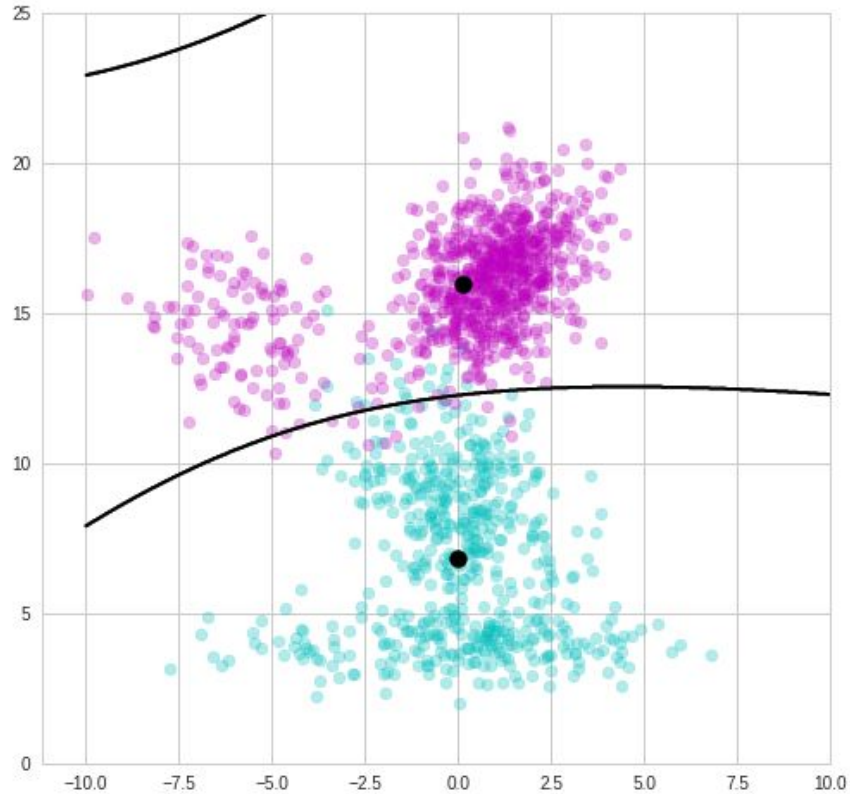
Gaussian Bayes Classifiers model the full covariance



naive training accuracy: 0.9286

bayes classifier training accuracy: 0.9657

Real data isn't as clean (which is why we get paid)



Creating mixture model Bayes classifiers is simple

```
gmm_a = GeneralMixtureModel.from_samples(MultivariateGaussianDistribution, 2, X[y == 0])  
gmm_b = GeneralMixtureModel.from_samples(MultivariateGaussianDistribution, 2, X[y == 1])  
model_b = BayesClassifier([gmm_a, gmm_b], weights=numpy.array([1-y.mean(), y.mean()]))
```

Creating any Bayes classifiers is simple

```
mc_a = MarkovChain.from_samples(X[y == 0])
mc_b = MarkovChain.from_samples(X[y == 1])
model_b = BayesClassifier([mc_a, mc_b], weights=numpy.array([1-y.mean(), y.mean()]))
```

```
hmm_a = HiddenMarkovModel...
hmm_b = HiddenMarkovModel...
model_b = BayesClassifier([hmm_a, hmm_b], weights=numpy.array([1-y.mean(), y.mean()]))
```

```
bn_a = BayesianNetwork.from_samples(X[y == 0])
bn_b = BayesianNetwork.from_samples(X[y == 1])
model_b = BayesClassifier([bn_a, bn_b], weights=numpy.array([1-y.mean(), y.mean()]))
```

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Finale: Train a mixture of hidden markov models in parallel

pomegranate has built in parallelization

```
%timeit model.predict(X)
%timeit predict(model, X, n_jobs=1)
%timeit predict(model, X, n_jobs=4)
```

```
1 loop, best of 3: 3.79 s per loop
1 loop, best of 3: 3.78 s per loop
1 loop, best of 3: 2.05 s per loop
```

```
print "Complete Match: ", (model.predict(X) == predict(model, X, n_jobs=4)).all()
print model.predict(X[:20])
print predict(model, X[:20], n_jobs=4)
```

```
Complete Match:  True
```

```
[0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0]
[0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0]
```

pomegranate has built in parallelization

```
model = NaiveBayes(distributions)
```

```
%timeit model.predict_proba(X)
```

```
%timeit predict_proba(model, X, n_jobs=4)
```

```
1 loop, best of 3: 6.43 s per loop
```

```
1 loop, best of 3: 3.53 s per loop
```

```
(model.predict_proba(X[:100]) - predict_proba(model, X[:100], n_jobs=4)).sum()
```

```
0.0
```


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Mixture of Hidden Markov Models

Creating a mixture of HMMs is just as simple as passing the HMMs into a GMM as if it were any other distribution

```
model_C = create_profile_hmm(dC, I)
model_mC = create_profile_hmm(dmC, I)
model_hmC = create_profile_hmm(dhmC, I)

model = GeneralMixtureModel([model_C, model_mC, model_hmC])
return model
```

parallel training of a mixture of hmms

Creation is just as simple as passing the HMMs into the GMM object. In this case, each model has 307 edges and 39 states to train

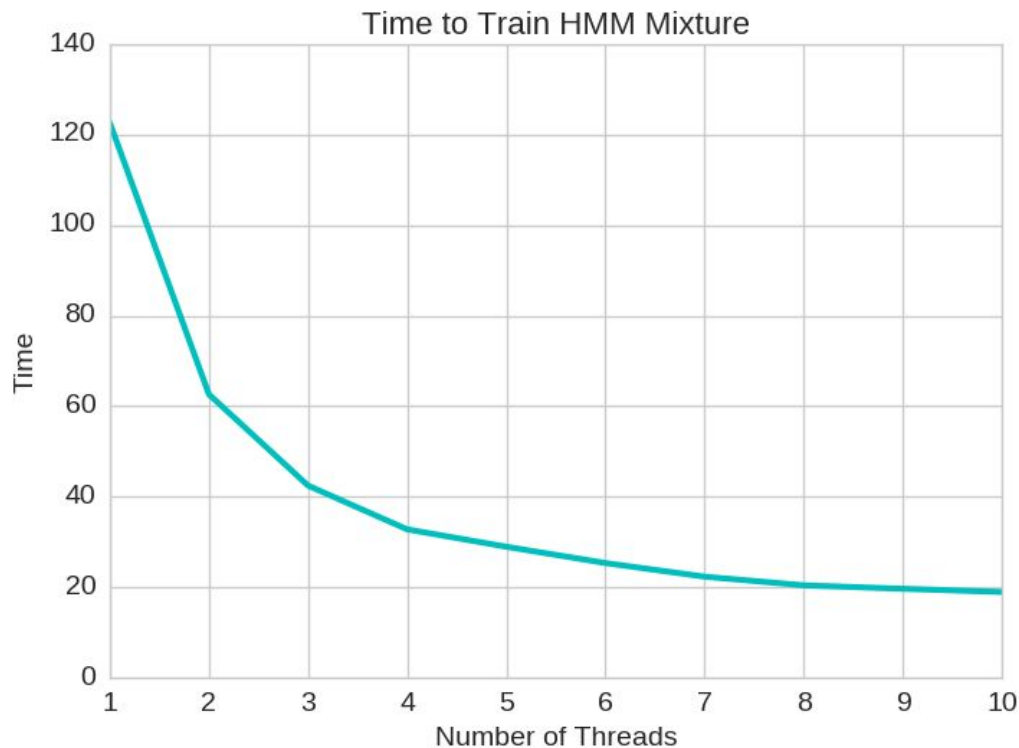
```
model_C = create_profile_hmm(dC, I)
model_mC = create_profile_hmm(dmC, I)
model_hmC = create_profile_hmm(dhmC, I)

model = GeneralMixtureModel([model_C, model_mC, model_hmC])
return model
```

```
model = create_mixture_model()
model.predict_proba([model.distributions[0].sample(), model.distributions[1].sample(), model.distributions[2].sample()])

array([[ 0.99999576,  0.00000413,  0.00000011],
       [ 0.00084806,  0.99915193,  0.00000001],
       [ 0.00001259,  0.00000395,  0.99998346]])
```

Parallel Training of a Mixture of HMMs



`fit(model, X, n_jobs=n)`

Overview

pomegranate can do **more** than other packages, **faster**, is **intuitive to use**, and can do it in **parallel**

Tutorials for each model are available on github

Branch: master ▾


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<https://github.com/jmschrei/pomegranate/tree/master/tutorials>

Thank you for your time.